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OM protein - protein search, using sw model

Run on: April 23, 2004, 16:21:29 / Search time 57 seconds  
(without alignments)  
736.589 Million cell updates/sec

Title: US-09-931-325c-170\_COPY\_1\_149  
Perfect score: 793  
Sequence: 1 MDIDYKFGATVLLSLP.....PAYRPPNAPILSTLPETTVV 149

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1586107 seqs, 282547505 residues

Total number of hits satisfying chosen parameters: 1586107

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : A\_Geneseq\_29Jan04: \*  
1: Geneseq1980s: \*  
2: Geneseq1990s: \*  
3: Geneseq2000s: \*  
4: Geneseq2001s: \*  
5: Geneseq2002s: \*  
6: Geneseq2003as: \*  
7: Geneseq2003bs: \*  
8: Geneseq2004s: \*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	793	100.0	171	5	AAU93972	Immunogen
2	793	100.0	183	2	AAV29674	Human hep
3	793	100.0	183	5	AAU87796	Human hep
4	793	100.0	183	5	AAU80906	Hepatitis
5	793	100.0	183	5	ABG94182	Hepatitis
6	793	100.0	183	5	AAU93961	Hepatitis
7	793	100.0	183	5	ABG80494	Hepatitis
8	793	100.0	183	6	ABO01919	Hepatitis
9	793	100.0	183	6	ABR56458	Hepatitis
10	793	100.0	183	6	ABR44571	Hepatitis
11	793	100.0	183	7	ADD24146	Hepatitis
12	793	100.0	183	7	AAE10767	Human Hep
13	793	100.0	183	2	AAW50242	Hepatitis
14	793	100.0	212	2	AAW50250	Hepatitis
15	793	100.0	212	5	AAU80921	Hepatitis
16	793	100.0	212	5	ABG94187	Hepatitis
17	793	100.0	212	5	AAE19898	Hepatitis
18	793	100.0	212	5	AAE19793	Hepatitis
19	793	100.0	212	5	ABG80509	Hepatitis
20	793	100.0	212	6	ABR56483	Hepatitis
21	793	100.0	212	6	ABR56483	Hepatitis
22	793	100.0	212	6	ABR44586	Hepatitis
23	793	100.0	212	7	ABW00349	Hepatitis
24	793	100.0	212	7	ADD24161	Hepatitis
25	793	100.0	289	2	AAW09049	Plasmid P

26	793	100.0	346	2	AAU27473	SI2/core
27	793	100.0	397	2	AAW09048	Plasmid P
28	791	99.7	183	5	AAU80909	Hepatitis
29	791	99.7	183	5	ABG94185	Hepatitis
30	791	99.7	183	5	ABG80497	Hepatitis
31	791	99.7	183	6	ABR56471	Synthetic
32	791	99.7	183	6	ABR44574	Synthetic
33	791	99.7	183	7	ADD24149	Hepatitis
34	790	99.6	183	5	AAU80911	Hepatitis
35	790	99.6	183	5	AAU80912	Hepatitis
36	790	99.6	183	5	ABG94187	Hepatitis
37	790	99.6	183	5	ABG94188	Hepatitis
38	790	99.6	183	5	ABG80500	Hepatitis
39	790	99.6	183	5	ABG80499	Hepatitis
40	790	99.6	183	6	ABR56473	Hepatitis
41	790	99.6	183	6	ABR56474	Hepatitis
42	790	99.6	183	6	ABR44576	Hepatitis
43	790	99.6	183	6	ABR44577	Hepatitis
44	790	99.6	183	7	ADD24151	Hepatitis
45	790	99.6	183	7	ADD24152	Hepatitis

ALIGNMENTS

RESULT 1

AAU93972  
ID AAU93972 standard; peptide; 171 AA.  
XX  
AC AAU93972;  
XX  
DT 02-JUL-2002 (first entry)  
XX  
DE Immunogenic HBC chimeric particle #9.  
XX  
KW Immunogenic; hepatitis nucleocapsid protein; hepatitis B core; HBC;  
KW vaccine; B cell epitope; T cell epitope; immunostimulant.  
XX  
OS Plasmodium falciparum.  
XX  
FN WO200214478-A2.  
XX  
PD 21-FEB-2002.  
XX  
PF 16-AUG-2001; 2001WO-US041759.  
XX  
PR 16-AUG-2000; 2000US-0225843P.  
PR 22-AUG-2000; 2000US-0226867P.  
PR 15-AUG-2001; 2001US-00930915.  
XX (APOV-) APOVIA INC.  
XX Birkett AJ;  
XX  
XX WPI; 2002-257601/30.  
XX  
XX Novel recombinant hepatitis B core protein, termed as chimeric  
XX hepatitis B core protein, displays immunogenic epitopes at N-terminus,  
XX HBC immunogenic loop with linker for conjugated epitope and C-terminus.  
XX  
XX Example 4; Page 273; 289pp; English.

The invention relates to a recombinant hepatitis nucleocapsid protein, i.e. a chimeric hepatitis B core (HBC) protein (I), displaying one or more immunogenic epitopes at the N-terminus, HBC immunogenic loop (L) or C-terminus, or having a heterologous linker for a conjugated epitope in (L), and containing a Cys residue at, or near, the C-terminus that confers enhanced stability to the particles. A vaccine comprising (I) is useful for inducing an immune response in an inoculated host animal, by inoculating a host animal with the vaccine, and maintaining that inoculated animal for a time period sufficient for that animal to develop an immune response. The immunogenic particles formed using (I) are substantially free of binding to nucleic acids, and are most stable than

CC the particle formed from otherwise identical HBC chimera that lacks the C-  
CC terminal residue or in which a C-terminal Cys is replaced by another  
CC residue. The chimera particles are most stable on storage in aqueous  
CC compositions that are particles of similar sequence that lack any C-  
CC terminal Cys residues. The chimera molecule exhibits the self-assembly not  
CC exhibiting the nucleic acid binding of those native particles, and  
CC excellent B cell and T cell immunogenicities. The chimera particles are  
CC typically prepared in higher yield than similar particles that are free  
CC of a C-terminal Cys. The particles are often far more immunogenic than  
CC the similar conjugates that lack a C-terminal Cys. Immunogenicities of  
CC particles assembled from the chimera molecules are enhanced as compared to  
CC similar particles assembled from chimera molecules lacking at least one C-  
CC terminal Cys. AAU93802-AAU93997 represent immunogenic HBC particles amino  
CC acid sequences and related sequences of the invention  
XX  
SQ Sequence 171 AA;

Query Match 100.0%; Score 793; DB 5; Length 171;  
Best Local Similarity 100.0%; Pred. No. 8.8e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKPRQLLWTHFHSCLTFTGRTVIEYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKPRQLLWTHFHSCLTFTGRTVIEYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 2  
AAU93674  
ID AAU93674 standard; protein; 183 AA.  
XX  
AC AAU93674;  
DT 08-NOV-1999 (first entry)  
XX  
DE Human hepatitis B core protein.  
XX  
KW Human hepatitis B core protein; HBC; modified; immunodominant;  
KW nucleocapsid protein; vaccine; T cell epitope.  
XX  
OS Hepatitis B virus.  
XX  
PN WO9940934-A1.  
PD 19-AUG-1999.  
XX  
PF 11-FEB-1999; 99WO-US003055.  
XX  
PR 12-FEB-1998; 98US-0074537P.  
XX  
PA (IMMU-) IMMUNE COMPLEX CORP.  
XX  
PI Birkett AJ;  
XX  
PI WPI; 1999-527340/44.  
DR N-PSDB; AAZ08816.  
XX

Conjugate of hepatitis B core protein, modified to increase reactivity  
PT with haptens, used to raise antibodies against the haptens, e.g. in  
PT vaccines.  
XX  
PS Claim 17; Page 77-78; 128pp; English.  
XX

The present invention describes a conjugate (A) comprising a  
CC strategically modified hepatitis B core (HBC) protein (I) attached to a  
CC haptens, where (I) includes amino acids (aa) 10-140 of the wild type HBC

CC 183 aa sequence (given in AAU92674) and additionally has an insert (II)  
CC in the region corresponding to aa's 50-100, where the insert is of 1 to  
CC about 40 aa's and contains a chemically reactive aa residue linked to the  
CC haptens. A vaccine containing (A), optionally in the form of particles, is  
CC used to induce a protective antibody response against the pathogen from  
CC which the haptens is derived, in humans or other animals. These pathogens  
CC may be bacteria, viruses, rickettsia or protozoa. Insertion of (II)  
CC overcomes the low reactivity of aa side chains in native HBC protein,  
CC increasing the reactivity with haptens and resulting in conjugates of  
CC improved immunogenicity. Modified HBC can be derivatised in the form of  
CC particles by well-defined chemical methods, and is unlikely to cause  
CC immunological side-effects. The present sequence represents the wild type  
CC HBC protein  
XX  
SQ Sequence 183 AA;

Query Match 100.0%; Score 793; DB 2; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKPRQLLWTHFHSCLTFTGRTVIEYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKPRQLLWTHFHSCLTFTGRTVIEYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 3  
AAU87796  
ID AAU87796 standard; protein; 183 AA.  
XX  
AC AAU87796;  
DT 21-MAY-2002 (first entry)  
XX  
DE Human hepatitis B virus nucleocapsid protein #1.  
XX  
KW Hepatitis B virus; nucleocapsid protein; HBC; hepatitis B virus core;  
KW B cell epitope; T cell epitope; malaria; HIV-1; Plasmodium; protozoa;  
KW circumsporozoite; human immunodeficiency virus type I; human; squirrel;  
KW woodchuck.  
XX  
OS Homo sapiens.  
XX  
PN WO200213765-A2.  
PD 21-FEB-2002.  
XX  
PF 16-AUG-2001; 2001WO-US025625.  
XX  
PR 16-AUG-2000; 2000US-0225813P.  
PR 15-AUG-2001; 2001US-00931325.  
XX  
PA (APOV-) APOVIA INC.  
XX  
PI Birkett AJ;  
XX  
PI WPI; 2002-241832/29.  
DR N-PSDB; ABK44278.  
XX

Recombinant hepatitis B virus core protein chimera molecule, useful to  
PT induce antibodies to malarial parasites, comprises malaria-specific T-  
PT cell epitope and is engineered for enhanced stability.  
XX  
PS Disclosure; Fig 6; 197pp; English.  
XX  
XX The invention relates to a recombinant hepatitis B virus core (HBC)

CC protein chimera molecule that contains 4 peptide-linked amino acid residue  
 CC sequence domains. The molecule of the invention contains a region  
 CC constituting a B cell epitope of the circumsporozoite protein of a  
 CC species of the parasite, Plasmodium. The chimera sequence is useful as an  
 CC immunogen for inducing antibodies to the malaria-causing parasite.  
 CC Plasmodium, particularly P. falciparum and P. vivax. Sequences AAU87695-  
 CC AAU87804 represent peptide epitopes of the invention  
 XX  
 XX Sequence 183 AA;

Query Match 100.0%; Score 793; DB 5; Length 183;  
 Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
 Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRTVIEYL 120  
 DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRTVIEYL 120  
 QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
 DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 4  
 AAU80906  
 ID AAU80906 standard; protein; 183 AA.  
 AC AAU80906;  
 XX  
 XX 09-APR-2002 (first entry)  
 DE Hepatitis B virus core antigen variant (HBCAg) #11.  
 DE Vaccine; molecular scaffold; pilus; pilin; HBCAg; antigen;  
 KW hepatitis B virus capsid protein; JUN; FOS; HIV Gp140;  
 KW measles virus N protein; bee venom phospholipase; Th type 2 T-helper;  
 KW Th2; Sinbis virus E2 protein; amyloid beta; influenza M2 antigen;  
 KW human immunodeficiency virus infection; viral hepatitis; measles;  
 KW chicken pox; pneumonia; tuberculosis; syphilis; malaria; allergy; cancer;  
 KW chronic disease; arthritis; colitis; diabetes; multiple sclerosis.  
 XX  
 XX Hepatitis B virus.  
 XX WO200185208-A2.  
 XX 15-NOV-2001.  
 XX 02-MAY-2001; 2001WO-IB000741.  
 XX 05-MAY-2000; 2000US-0202341P.  
 XX (CYTO-) CYTOS BIOTECHNOLOGY AG.  
 XX (SEBB) SEBBEL P.  
 XX (DUNA/) DUNANT N.  
 XX (BACH/) BACHMANN M.  
 XX (TISS/) TISSOT A.  
 XX (LECH/) LECHENER F.  
 XX Sebbel P, Dunant N, Bachmann M, Tissot A, Lechener F;  
 PI WPI; 2002-055561/07.  
 DR  
 XX New composition, useful for vaccine production, comprises antigen or  
 PT antigenic determinant and non-natural molecular scaffold comprising  
 PT organizer and core particle such as bacterial pilus or pilin protein.  
 XX  
 PS Claim 34; Page 216-217; 287pp; English.  
 XX The invention relates to a composition comprising: (a) a non-natural

CC molecular scaffold (molecular scaffold) which comprises a core particle  
 CC such as a bacterial pilus or pilin protein, a recombinant form of the  
 CC protein, a virus-like particle or a hepatitis B virus capsid protein  
 CC (HBCAg), and an organizer; and (b) an antigen or antigenic determinant,  
 CC where the molecular scaffold and antigenic determinant interact to form  
 CC an ordered and repetitive antigen array. Suitable antigenic determinants  
 CC include JUN, FOS, HIV Gp140, measles virus N protein, bee venom  
 CC phospholipase, Sinbis virus E2 protein, amyloid beta derived peptides and  
 CC influenza M2 antigen. The composition (or vaccine) is useful for  
 CC immunisation, by administration to a subject, where the administration  
 CC produces an immune response, such as humoral, cellular or protective  
 CC immune response, preferably a Th type 2 T-helper (Th2) response that is  
 CC specific for the antigenic determinant. The administration induces  
 CC antibodies specific for the antigenic determinant of a subtype  
 CC corresponding to the Th2 subtype in the subject. The subject does not  
 CC generate a Th2 subtype that is specific for pilus or pilin polypeptide or  
 CC antigenic determinant. The composition is useful for the production of  
 CC vaccines for prevention of infectious diseases such as human  
 CC immunodeficiency virus, viral hepatitis, measles, chicken pox, pneumonia,  
 CC tuberculosis, syphilis, malaria, and for treating allergy, cancer, and  
 CC chronic diseases induced or accelerated by a Th1 type immune response,  
 CC such as arthritis, colitis, diabetes and multiple sclerosis. The  
 CC composition is useful to generate defined self-specific antibodies and  
 CC specific immune responses of the Th2 type and allows the creation of  
 CC highly efficient vaccines against infectious diseases, and for treating  
 CC allergy, cancer, and chronic diseases induced or accelerated by a Th1  
 CC type immune response. The present sequence is a peptide or protein  
 CC incorporated into the compositions of the invention  
 XX  
 XX Sequence 183 AA;

Query Match 100.0%; Score 793; DB 5; Length 183;  
 Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
 Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRTVIEYL 120  
 DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRTVIEYL 120  
 QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
 DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 5  
 ABG94182  
 ID ABG94182 standard; protein; 183 AA.  
 XX  
 XX ABG94182;  
 XX  
 XX 10-DEC-2002 (first entry)  
 DT Hepatitis B capsid (core) protein antigen (HBCAg) variant #11.  
 XX Human; mouse; rat; antimicrobial; antiallergic; immunomodulatory;  
 XX cytoskeletal; antiviral; antidiabetic; hypoglycaemic; antigen array;  
 KW vaccine; infectious disease.  
 XX  
 XX Hepatitis B virus.  
 OS  
 XX WO200256905-A2.  
 XX  
 XX 25-JUL-2002.  
 XX  
 XX 21-JAN-2002; 2002WO-IB000166.  
 PF  
 XX 19-JAN-2001; 2001US-0262379P.  
 PR 04-MAY-2001; 2001US-0288549P.  
 PR 05-OCT-2001; 2001US-0326998P.

PR 07-NOV-2001; 2001US-0331045P.  
XX (CYTO-) CYTOS BIOTECHNOLOGY AG.  
XX Renner WA, Bachmann M, Tissot A, Maurer P, Lechner F, Sebbel P;  
PI Piossek C;  
XX WPI; 2002-627351/67.  
XX Molecular antigen array used in the production of vaccines for infectious  
PT diseases.  
XX Claim 112; Page 337-338; 441pp; English.  
XX This invention relates to a novel ordered and repetitive antigen array  
CC used in the production of vaccines for infectious diseases. The invention  
CC also discloses a composition comprising a non-natural molecular scaffold  
CC comprising a core particle selected from a core particle of a non-natural  
CC origin and a core particle of natural origin and an organiser comprising  
CC at least one first attachment site, where the organiser is connected to  
CC the core particle by at least one covalent bond. Also disclosed is an  
CC antigen or antigenic determinant with at least one second attachment  
CC site, where the antigen or antigenic determinant is anyloid beta peptide  
CC (Abeta1-42) or its fragment and where the second attachment site is  
CC selected from an attachment site not naturally occurring with the antigen  
CC or antigenic determinant and an attachment site naturally occurring with  
CC the antigen or antigenic determinant, where the second attachment site is  
CC capable of association through at least one non-peptide bond to the first  
CC attachment site and where the antigen or antigenic determinant and the  
CC scaffold interact through the association to form an ordered and  
CC repetitive antigen array. The invention also comprises a coat protein  
CC capable of forming a capsid which comprises mutant beta coat proteins  
CC having an amino acid sequence selected from five amino acid sequences  
CC fully defined in the specification. The compounds of the invention may  
CC have antimicrobial, anti-allergic, immunomodulatory, cytostatic,  
CC antiviral, antidiabetic, or hypoglycaemic activities and may be used in  
CC immunisation and as a vaccine. The present sequence represents a protein  
XX sequence used to create the compositions of the invention  
XX  
SQ Sequence 183 AA;  
Query Match 100.0%; Score 793; DB 5; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALQAIL 60  
Db 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALQAIL 60  
Qy 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEYLV 120  
Db 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEYLV 120  
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
RESULT 6  
AAU93961  
ID AAU93961 standard; peptide; 183 AA.  
XX AAU93961;  
XX  
XX 02-JUL-2002 (first entry)  
XX Hepatitis B virus Hbc epitope #1.  
XX Immunogenic; hepatitis B virus nucleocapsid protein; hepatitis B core; Hbc;  
XX vaccine; B cell epitope; T cell epitope; immunostimulant.  
XX  
XX Hepatitis B virus.  
XX

PN WO200214478-A2.  
XX  
PD 21-FEB-2002.  
XX  
EP 16-AUG-2001; 2001WO-US041759.  
XX  
XX 16-AUG-2000; 2000US-0225843P.  
PR 22-AUG-2000; 2000US-0226867P.  
PR 15-AUG-2001; 2001US-00930915.  
XX  
XX (APOV-) APOVIA INC.  
XX  
XX Birkett AJ;  
XX WPI; 2002-257601/30.  
XX Novel recombinant hepatitis B core protein, displaying one or  
PT hepatitis B core protein, displays immunogenic epitopes at N-terminus,  
PT Hbc immunogenic loop with linker for conjugated epitope and C-terminus.  
XX  
XX Disclosure; Fig 7; 289pp; English.  
XX The invention relates to a recombinant hepatitis B core protein, termed as chimeric  
CC i.e. a chimeric hepatitis B core (Hbc) protein (I), displaying one or  
CC more immunogenic epitopes at the N-terminus, Hbc immunogenic loop (L) or  
CC C-terminus, or having a heterologous linker for a conjugated epitope in  
CC (L), and containing a Cys residue at, or near, the C-terminus that  
CC confers enhanced stability to the particles. A vaccine comprising (I) is  
CC useful for inducing an immune response in an inoculated host animal, by  
CC inoculating a host animal with the vaccine, and maintaining that  
CC inoculated animal for a time period sufficient for that animal to develop  
CC an immune response. The immunogenic particles formed using (I) are  
CC substantially free of binding to nucleic acids, and are most stable than  
CC the particle formed from otherwise identical Hbc chimera that lacks the C-  
CC terminal residue or in which a C-terminal Cys is replaced by another  
CC residue. The chimera particles are most stable on storage in aqueous  
CC compositions that are particles of similar sequence that lack any C-  
CC terminal Cys residues. The chimera molecule exhibits the self-assembly not  
CC exhibiting the nucleic acid binding of those native particles, and  
CC excellent B cell and T cell immunogenicities. The chimera particles are  
CC typically prepared in higher yield than similar particles that are free  
CC of a C-terminal Cys. The particles are often far more immunogenic than  
CC the similar conjugates that lack a C-terminal Cys. Immunogenicities of  
CC similar particles assembled from the chimera molecules are enhanced as compared to  
CC similar particles assembled from chimera molecules lacking at least one C-  
CC terminal Cys. AAU93961-AAU93997 represent immunogenic Hbc particles amino  
CC acid sequences and related sequences of the invention  
XX  
SQ Sequence 183 AA;  
Query Match 100.0%; Score 793; DB 5; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
Qy 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALQAIL 60  
Db 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALQAIL 60  
Qy 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEYLV 120  
Db 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEYLV 120  
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
RESULT 7  
ABG80494  
ID ABG80494 standard; protein; 183 AA.  
XX  
XX ABG80494;  
XX

DT XX 29-NOV-2002 (first entry)  
DE XX Hepatitis B virus core capsid protein, HBcAg, variant #11.  
KW Molecular antigen array; vaccine; antigen; antimicrobial;  
KW molecular scaffold; amyloid beta; Abeta 1-42; influenza;  
KW graft versus host disease; Igs-mediated allergic reaction; anaphylaxis;  
KW adult respiratory distress syndrome; ARDS; Crohn's disease;  
KW allergic asthma; acute lymphoblastic leukaemia; non-Hodgkin's lymphoma;  
KW Grave's disease; systemic lupus erythematosus; osteoporosis;  
KW inflammatory immune disease; myasthenia gravis; multiple sclerosis;  
KW immunoproliferative disease lymphadenopathy; Alzheimer's disease;  
KW angioimmunoproliferative lymphadenopathy; immunoblastic lymphadenopathy;  
KW rheumatoid arthritis; diabetes; infectious disease; factor Xa;  
KW enterokinase; cysteine-containing linker.  
XX OS Hepatitis B virus.  
XX XX WO200256907-A2.  
XX PD 25-JUL-2002.  
XX XX 21-JAN-2002; 2002WO-IB000168.  
XX XX 19-JAN-2001; 2001US-0262379P.  
XX PR 04-MAY-2001; 2001US-0286549P.  
XX PR 05-OCT-2001; 2001US-0326998P.  
XX PR 07-NOV-2001; 2001US-0331045P.  
XX XX (CYTO-) CYTOS BIOTECHNOLOGY AG.  
XX PA (NOVS) NOVARTIS PHARMA AG.  
XX PA (MAUR) MAURER P.  
XX PA (LECH) LECHNER F.  
XX PA (ORTM) ORTMANN R.  
XX PA (LUBO) LUBOEND R.  
XX PA (STAU) STAUFENBIEL M.  
XX PA (FREY) FREY P.  
XX XX Maurer P, Lechner F, Ortmann R, Lueoend R, Staufenbiel M, Frey P;  
PI Renner WA, Bachmann M, Tissot A, Sebbel P, Piossek C;  
XX WPI; 2002-636514/68.  
XX DR Molecular antigen array used in the production of vaccines for infectious  
XX PT diseases.  
XX PS Claim 14; Page 314-315; 418pp; English.  
XX CC The invention relates to a composition comprising: (a) a non-natural  
XX molecular scaffold comprising: (i) a core particle selected from: (1) a  
XX core particle of a non-natural origin; and (2) a core particle of natural  
XX origin; and (ii) an organiser comprising at least one first attachment  
XX site, where the organiser is connected to the core particle by at least  
XX one covalent bond; (b) an antigen or antigenic determinant with at least  
XX one second attachment site, where the antigen or antigenic determinant is  
XX amyloid beta peptide (Abeta 1-42) or its fragment, and where the second  
XX attachment site is selected from: (i) an attachment site not naturally  
XX occurring with the antigen or antigenic determinant; and (ii) an  
XX attachment site naturally occurring with the antigen or antigenic  
XX determinant, where the second attachment site is capable of association  
XX through at least one non-peptide bond to the first attachment site; and  
XX where the antigen or antigenic determinant and the scaffold interact  
XX through the association to form an ordered and repetitive antigen array.  
XX Also included is a process for producing a non-naturally occurring  
XX ordered and repetitive antigen array. The composition is used in  
XX immunisation and as a vaccine for diseases such as influenza, graft  
XX versus host disease, Igs-mediated allergic reactions, anaphylaxis, adult  
XX respiratory distress syndrome (ARDS), Crohn's disease, allergic asthma,  
XX acute lymphoblastic leukaemia, non-Hodgkin's lymphoma, Grave's disease,  
XX systemic lupus erythematosus, inflammatory immune diseases, myasthenia  
XX gravis, immunoproliferative disease lymphadenopathy,  
XX angioimmunoproliferative lymphadenopathy, immunoblastic lymphadenopathy,  
XX rheumatoid arthritis, diabetes, multiple sclerosis, Alzheimer's disease,  
XX

CC osteoporosis and infectious diseases. The present sequence is an antigen  
CC for use in the array of the invention. The antigen is modified to possess  
CC a cleavage site (enterokinase or factor Xa) and a Cysteine- containing N-  
CC or C-terminal linker peptide which serves as the attachment point to a  
CC virus like particle or bacterial protein (the scaffold protein)  
XX SQ Sequence 183 AA;  
XX Query Match 100.0%; Score 793; DB 5; Length 183;  
XX Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
XX Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPKFEGATVELLSFLPSPDFPSVRLDITASALYREALSEPHCSPHHTALRQAIL 60  
DB 1 MDIDPKFEGATVELLSFLPSPDFPSVRLDITASALYREALSEPHCSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGLKFRQLLMFHFISCLTFGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGLKFRQLLMFHFISCLTFGRETVEIYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
XX RESULT 8  
XX ID ABO01919 standard; protein; 183 AA.  
XX AC ABO01919;  
XX XX 07-AUG-2003 (first entry)  
XX DT Hepatitis B virus core protein antigen.  
XX DE Cholesteryl ester transfer protein; CETP; HDL cholesterol;  
XX KW high density lipoprotein; immunogen; pro-atherogenic dyslipoproteinaemia;  
XX KW low density lipoprotein; LDL; HBcAg; hepatitis core antigen.  
XX OS Hepatitis B virus.  
XX XX US2003021804-A1.  
XX PD 30-JAN-2003.  
XX XX 21-JAN-1997; 97US-00785997.  
XX XX 21-JAN-1997; 97US-00785997.  
XX XX (NEED/) NEEDLEMAN P.  
XX PA (GLEN/) GLENN K.  
XX PI Needleman P, Glenn K;  
XX DR WPI; 2003-456282/43.  
XX DR N-PSDB; ACD07806.  
XX XX Increasing high density lipoprotein cholesterol level in blood of a  
XX mammal whose blood contains cholesteryl ester transfer protein, by  
XX PT immunizing mammal with inoculum of cholesteryl ester transfer protein  
XX PT immunogen.  
XX PS Disclosure; Page 30; 35pp; English.  
XX XX The invention relates to increasing high density lipoprotein (HDL)  
XX cholesterol in the blood of a mammal whose blood contains cholesteryl  
XX ester transfer protein (CETP), comprising immunising mammal with an  
XX inoculum having CETP immunogen that is an immunogenic polypeptide (IP)  
XX having CETP amino acid residue sequence, allowing IP to induce production  
XX of anti-CETP antibodies in mammal and lessen transfer of cholesteryl  
XX esters from HDL. Also included is a CETP immunogen that comprises an  
XX immunogenic polypeptide having a CETP amino acid residue sequence  
XX covalently bonded to an exogenous antigenic carrier (e.g. hepatitis B  
XX

core protein antigen, HBcAg). The method is useful for increasing the concentration of HDL cholesterol in the blood of a mammal whose blood contains CETP. The method is useful for ameliorating or treating pro-atherogenic dyslipoproteinaemias characterised by low HDL/low density lipoprotein (LDL) cholesterol ratios. The method has an effect that lasts for months as compared to the short-term effects of the small molecule drugs not available. The method utilises the host mammal's own (autogenic) immunological system to provide a desired result, thus obviating problems associated with repeated administration of xenogeneic antibodies that themselves become immunogenic in the host mammal. The method utilises accepted exogenous antigenic carriers such as hepatitis B core protein (HBcAg), tetanus toxoid, tuberculin purified protein derivative (PPD) and diphtheria toxoid which can boost the host mammal's immunity to those pathogens. The method also lessens the cholesterol ester transfer from HDL to LDL or very low density lipoprotein, thus increasing the concentration of anti-atherogenic HDL cholesterol. The present sequence represents the hepatitis B core protein (HBcAg) used as the exogenous antigenic carrier

Sequence 183 AA;

Query Match 100.0%; Score 793; DB 6; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 60  
DB 1 MDIDPKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTFGRVIEYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTFGRVIEYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

# RESULT 9

ID ABR56468 standard; protein; 183 AA.

XX ABR56468;  
AC ABR56468;

DT 28-JUL-2003 (first entry)

DE Hepatitis B core antigen precursor variant protein SEQ ID NO:39.

XX Antigen presenting cell; APC; immune response; virus like particle; VLP;  
KW cytostatic; virucide; antibacterial; antiparasitic; fungicide;  
KW antiallergic; immunosuppressive; antiaddictive; antiinflammatory;  
KW antithyroid; antidiabetic; neuroprotective; nootropic; osteopathic;  
KW antihumetic; antiarthritic; vaccine; immunisation; infectious disease;  
KW anti-viral protection; tumour; allergy; drug addiction; Crohn's disease;  
KW graft-versus-host disease; Grave's disease; diabetes; multiple sclerosis;  
KW Alzheimer's disease; osteoporosis; rheumatoid arthritis;  
KW inflammatory autoimmune disease.

XX Hepatitis B virus.  
OS Synthetic.

XX WO2003024480-A2.

XX 27-MAR-2003.

XX 16-SEP-2002; 2002WO-IB004252.

XX 14-SEP-2001; 2001US-0318967P.

XX (CYTO-) CYTOS BIOTECHNOLOGY AG.

XX Bachmann MF, Storni T, Lechner F;

DR MPI; 2003-363095/34.

XX A composition, useful for enhancing an immune response against an antigen or a virus-like particle, enhancing anti-viral protection in an animal, or immunizing or treating tumors or infectious diseases, e.g. viral infections.

PS Disclosure; Page 195; 243pp; English.

XX The present invention describes a composition (C) for enhancing an immune response against an antigen or a virus-like particle in an animal. (C) comprises a virus-like particle (VLP) bound to at least one antigen, or a VLP capable of being recognised by the immune system of the animal. Also described: (1) enhancing an immune response against an antigen or a VLP in an animal comprising introducing (C) into the animal; (2) vaccines comprising (C) together with a pharmaceutical diluent, carrier or excipient; (3) immunising or treating an animal comprising administering the vaccine to the animal, or priming or boosting a T cell response in the animal by administering the vaccine; and (4) enhancing anti-viral protection in an animal comprising introducing (C) into the animal. (C) has cytostatic, virucide, antibacterial, antiparasitic, fungicide, antiallergic, immunosuppressive, antiaddictive, antiinflammatory, antithyroid, antidiabetic, neuroprotective, nootropic, osteopathic, antihumetic and antiarthritic activities. (C) or the vaccines can be used for enhancing an immune response against an antigen or a VLP in an animal, enhancing anti-viral protection in an animal, or immunising or treating tumors and infectious diseases such as viral, bacterial, parasitic or fungal infections. The vaccine compositions are also useful for preventing or treating allergies, drug addiction, graft-versus-host disease, Crohn's disease, osteoporosis, rheumatoid arthritis, or inflammatory Alzheimer's disease. ACC69838 to ACC69852 and ABR56401 to ABR56509 represent sequences used in the exemplification of the present invention

XX Sequence 183 AA;

Query Match 100.0%; Score 793; DB 6; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9.7e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 60  
DB 1 MDIDPKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTFGRVIEYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTFGRVIEYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

# RESULT 10

ID ABR44571 standard; protein; 183 AA.

XX ABR44571;

XX 25-JUL-2003 (first entry)

DE Hepatitis B core antigen precursor variant protein SEQ ID NO:39.

XX Immunostimulatory; virus-like particle; bacteriophage; HBV; LCMV;  
KW hepatitis B virus; lymphocytic choriomeningitis virus; vaccine;  
KW immunostimulant; cytostatic; antiallergic; virucide; antibacterial;  
KW immune response; immunisation; allergy; tumour; breast cancer;  
KW neuroblastoma; leukaemia; viral disease; influenza; hepatitis; measles;  
KW chicken pox; bacterial infection; tuberculosis; pneumonia; syphilis.

XX Hepatitis B virus.

OS Synthetic.

XX

PN WO2003024481-A2.  
XX 27-MAR-2003.  
PD 16-SEP-2002; 2003WO-IB004132.  
XX 14-SEP-2001; 2001US-0318994P.  
XX 22-APR-2002; 2002US-0374145P.  
XX (CYTO-) CYTOS BIOTECHNOLOGY AG.  
PA (MAUR-) MAURER P.  
PA (TISS-) TISSOT A.  
PA (SCHW-) SCHWARZ K.  
PA (MEIJ-) MEIJERINK E.  
PA (LIPO-) LIPOWSKY G.  
PA (PUMP-) PUMPKENS P.  
PA (CIEL-) CIELENS I.  
PA (RENH-) RENHOFA R.  
XX Maurer P, Tissot A, Schwarz K, Meijerink E, Lipowsky G;  
PI Pumpens P, Cielens I, Renhofs R, Bachmann MP, Storni T;  
XX WPI; 2003-354564/33.  
DR New compositions comprising immunostimulatory substances packaged into  
XX virus-like particles, useful as a vaccine for enhancing an immune  
PT response in animals, e.g. for treating or preventing allergies, tumors or  
PT viral infections.  
XX Disclosure; Page 274-275; 322pp; English.  
XX The present invention describes a composition (C) for enhancing an immune  
CC response in an animal. (C) comprises a virus-like particle (VLP), and an  
CC immunostimulatory substance. The immunostimulatory substance is bound to  
CC the VLP. Also described: (1) enhancing an immune response in an animal by  
CC introducing (C) into the animal; (2) producing (C) for enhancing an  
CC immune response in an animal; (3) vaccines comprising (C) together with a  
CC pharmaceutical diluent, carrier or excipient; and (4) immunising or  
CC treating an animal by: (a) administering the vaccine to the animal; (b)  
CC priming a T cell response in the animal by administering the vaccine; or  
CC (c) boosting a T cell response in the animal by administering the  
CC vaccine. (C) has immunostimulant, cytostatic, antiallergic, virucide and  
CC antibacterial activities. (1) can be used in vaccines for enhancing an  
CC immune response in an animal, particularly a mammal or human.  
CC Specifically, (C) is useful for enhancing a B cell response, a T cell  
CC response, or a cytotoxic T-lymphocyte (CTL) response. (C) or a vaccine  
CC comprising (C) can also be used for immunising or treating an animal,  
CC e.g. humans, sheep, horses, cattle, pigs, dogs, cats, rats, birds,  
CC reptiles or fish. (C) is particularly useful in prophylactic or  
CC therapeutic vaccines against allergies, tumours (e.g. breast cancers,  
CC neuroblastoma, or leukaemia), viral diseases (e.g. influenza, hepatitis,  
CC measles or chicken pox), or bacterial infections (e.g. tuberculosis,  
CC pneumonia or syphilis). ACC59790 to ACC69815 and ABR44502 to ABR44612  
CC represent sequences used in the exemplification of the present invention  
XX  
XX Sequence 183 AA;  
Query Match 100.0%; Score 793; DB 6; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9,7e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 60  
Db 1 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRLVSVYNTNMGKFRQLLWFIHISCLTFTGRTVIEYLV 120  
Db 61 CWGELMTLATWGVNLEDPASRLVSVYNTNMGKFRQLLWFIHISCLTFTGRTVIEYLV 120  
QY 121 SFGWIRTTPAYRPPNAPILSTLPETTV 149  
Db 121 SFGWIRTTPAYRPPNAPILSTLPETTV 149

RESULT 11  
ADD24146  
ID ADD24146 standard; protein; 183 AA.  
XX AC ADD24146;  
XX DT 15-JAN-2004 (first entry)  
XX DE Hepatitis B virus core antigen (HBcAg) precursor variant Seq ID39.  
XX KW vaccine composition; virus-like particle; core particle;  
KW first attachment site; antigen; antigenic determinant; prion protein;  
KW PrP; PrP peptide; vaccine; neuroprotective; antiinflammatory;  
KW prion disease; Bovine Spongiform Encephalopathy; BSE;  
KW Creutzfeldt-Jakob Disease; HBcAg; mutant; muten.  
XX OS Hepatitis B virus.  
XX PN WO2003059386-A2.  
XX 24-JUL-2003.  
XX 17-JAN-2003; 2003WO-EP000460.  
XX 18-JAN-2002; 2002US-00050902.  
XX 21-JAN-2002; 2002WO-IB000166.  
XX 08-JUL-2002; 2002US-0393725P.  
XX 18-JUL-2002; 2002US-0396590P.  
XX (CYTO-) CYTOS BIOTECHNOLOGY AG.  
XX Bachmann M, Maurer P, Pelliccioli E, Renner WA;  
XX WPI; 2003-598493/56.  
XX A vaccine composition for preventing or treating prion diseases (e.g.  
PT Creutzfeldt-Jakob Disease) comprises a virus-like particle (e.g. RNA-  
PT phase) and at least one prion protein or peptide bound to the virus-like  
PT particle.  
XX Disclosure; SEQ ID NO 39; 246pp; English.  
XX This invention relates to a novel vaccine composition comprising a virus-  
XX like or a core particle with at least one first attachment site and at  
XX least one antigen or antigenic determinant that is a prion protein (PrP)  
XX or its dimer, or a PrP peptide, the antigen or antigenic determinant  
XX being bound to the virus-like or core particle. The vaccine of the  
XX invention may have neuroprotective or antiinflammatory activity. The  
XX composition is useful as a medicament or in manufacturing a medicament  
XX for the treatment or prevention of prion diseases. The prion diseases may  
XX include Bovine Spongiform Encephalopathy (BSE) or Creutzfeldt-Jakob  
XX Disease. The present sequence is the amino acid sequence of a mutant  
XX hepatitis B virus core antigen (HBcAg) which may be used during the  
XX creation of the vaccine composition of the invention.  
XX Sequence 183 AA;  
Query Match 100.0%; Score 793; DB 7; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9,7e-85;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 60  
Db 1 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRLVSVYNTNMGKFRQLLWFIHISCLTFTGRTVIEYLV 120  
Db 61 CWGELMTLATWGVNLEDPASRLVSVYNTNMGKFRQLLWFIHISCLTFTGRTVIEYLV 120  
QY 121 SFGWIRTTPAYRPPNAPILSTLPETTV 149  
Db 121 SFGWIRTTPAYRPPNAPILSTLPETTV 149

## RESULT 12

AD10767  
ID ADE10767 standard; protein; 183 AA.

AC ADE10767;  
AD10767;

DT 29-JAN-2004 (first entry)  
XX

DE Human Hepatitis B virus sub-type AYW core protein.  
XX

XX hepatotropic; virucide; antiinflammatory; chronic hepatitis; vaccine;  
KW recombinant hepatitis B core chimeric protein; Hbc chimeric protein;  
KW hepatitis B infection; T-cell stimulator.  
XX

OS Hepatitis B virus.  
XX

XX US2003198645-A1.  
PN

XX 23-OCT-2003.  
PD

XX 21-FEB-2003; 2003US-00372076.  
PF

XX 21-FEB-2002; 2002US-00080299.  
PR

XX 21-FEB-2002; 2002US-00082014.  
PR

XX (PAGE//) PAGE M.  
PA

XX (FRIE//) FRIEDE M.  
PA

XX Page M, Friede M;  
PI

XX WPI; 2003-852775/79.  
DR

XX N-PSDB; ADE10968.  
DR

XX Claim 18; SEQ ID NO 1; 111pp; English.  
PS

XX The invention describes a method of treating chronic hepatitis comprising  
CC administering to a patient a T cell-stimulating amount of a vaccine  
CC comprising immunogenic particles dissolved or dispersed in a diluent,  
CC where the immunogenic particles consists of recombinant hepatitis B core  
CC (Hbc) chimeric protein molecules, and maintaining the patient to induce T  
CC cells activated against Hbc. The methods and compositions of the present  
CC invention are useful for treating chronic hepatitis B infection. This is  
CC the amino acid sequence of a Hepatitis B core protein.  
XX

XX Sequence 183 AA;  
SQ

Query Match 100.0%; Score 793; DB 7; Length 183;  
Best Local Similarity 100.0%; Pred. No. 9.7e-85;

Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHTRQAIL 60

DB 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHTRQAIL 60

QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRETVEYLV 120

DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRETVEYLV 120

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

## RESULT 13

AAW50242  
ID AAW50242 standard; protein; 194 AA.

XX AAW50242;  
AC

DT 28-SEP-1998 (first entry)  
XX

DE Hepatitis B virus precore p22 polypeptide Met-p22.  
XX

XX Viral replication; inhibitor; HBV; nucleocapsid; gene therapy;  
KW hepatocyte; liver; Met-p22.  
XX

OS Hepatitis B virus.  
OS

OS Synthetic.  
OS

XX Key Location/Qualifiers  
FH Protein 2..194

XX /label= p22  
FT

XX WO9809649-A1.  
PN

XX 12-MAR-1998.  
PD

XX 03-SEP-1997; 97WO-US015500.  
PF

XX 03-SEP-1996; 96US-0025370P.  
PR

XX (GEHO) GEN HOSPITAL CORP.  
PA

XX Wands JR, Scaglioni PP, Melegari M;  
PI

XX WPI; 1998-193325/17.  
DR

XX DNA encoding proteins which can be incorporated with wild type  
PT nucleocapsid subunit(s) into a viral nucleocapsid - useful for  
PT inhibition of viral replication, especially hepatitis B virus.  
PT

XX Claim 11; Page 40; 60pp; English.  
PS

XX This polypeptide comprises the hepatitis B virus (HBV) 22 kDa (p22)  
CC protein with an added N-terminal Met residue. p22 is produced by  
CC elimination of the 19-amino acid leader peptide from the 25 kDa full-  
CC length HBV precore protein (see AAW50250). Evidence is provided that HBV  
CC replication is inhibited in the presence of high levels of HBV precore or  
CC precore-related proteins. These proteins can be incorporated into HBV  
CC nucleocapsids along with the p21 core protein (see AAW50251), which is  
CC the usual nucleocapsid component, and thereby render the nucleocapsids  
CC deficient in encapsidating HBV pregenomic RNA. Thus, over-expression of  
CC the precore proteins, or certain variants of them, leads to transdominant  
CC inhibition of HBV replication. Suitable inhibitory proteins include p25  
CC (see AAW50250), p22, Met-p22, p18 (see AAW50236), Met-p18 (see AAW50237)  
CC and Met-p18-Het (see AAW50238). Heterologous peptides (see AAW50244-49)  
CC may be inserted into the p22 and Met-p22 polypeptides. The inhibitory  
CC proteins can be produced by recombinant methods using claimed expression  
CC vectors and host cells. They can be provided exogenously to the target  
CC cells for use in inhibiting HBV replication. Alternatively, a nucleic  
CC acid construct that directs overexpression of an inhibitory protein in  
CC target cells is used for the gene therapy of HBV infection  
XX

XX Sequence 194 AA;  
SQ

Query Match 100.0%; Score 793; DB 2; Length 194;  
Best Local Similarity 100.0%; Pred. No. 1e-84;

Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHTRQAIL 60

DB 12 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHTRQAIL 71

QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRETVEYLV 120

DB 72 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWPHISCLTFGRETVEYLV 131

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149



Db 132 SFGWIRTPPAYRPPNAPILSTLPETTV 160

## RESULT 14

AAW50250  
ID AAW50250 standard; protein; 212 AA.

XX AC AAW50250;  
XX DT 28-SEP-1998 (first entry)

XX DE Hepatitis B virus precore p25 polypeptide.

XX KW Viral replication; inhibitor; HBV; nucleocapsid; gene therapy;  
XX KW hepatocyte; liver; p25 protein.  
XX OS Hepatitis B virus.  
XX PN WO9809649-A1.

XX PD 12-MAR-1998.

XX PF 03-SEP-1997; 97WO-US015500.

XX PR 03-SEP-1996; 96US-0025370P.

XX PA (GEO ) GEN HOSPITAL CORP.

XX PI Wands JR, Scaglioni PP, Melegari M;

XX WPI; 1998-193325/17.

XX DNA encoding proteins which can be incorporated with wild type  
XX PT nucleocapsid subunit(s) into a viral nucleocapsid - useful for  
XX PT inhibition of viral replication, especially hepatitis B virus.  
XX PS Claim 15; Page 35; 60pp; English.

XX This polypeptide comprises the hepatitis B virus (HBV) 25 kDa (p25)  
XX protein that is encoded by the full-length HBV precore gene. Evidence is  
XX provided that HBV replication is inhibited in the presence of high levels  
XX of HBV precore or precore-related proteins. These proteins can be  
XX incorporated into HBV nucleocapsids along with the p21 core protein (see  
XX AAW50251), which is the usual nucleocapsid component, and thereby render  
XX the nucleocapsids deficient in encapsidating HBV pregenomic RNA. Thus,  
XX over-expression of the precore proteins, or certain variants of them,  
XX leads to transdominant inhibition of HBV replication. Suitable inhibitory  
XX proteins include p25, p22 (see AAW50241), Met-p22 (see AAW50242), p18  
XX (see AAW50236), Met-p18 (see AAW50237) and Met-p18-Het (see AAW50238).  
XX The inhibitory proteins can be produced by recombinant methods using  
XX claimed expression vectors and host cells. They can be provided  
XX exogenously to the target cells for use in inhibiting HBV replication.  
XX Alternatively, a nucleic acid construct that directs overexpression of an  
XX inhibitory protein in target cells is used for the gene therapy of HBV  
XX infection.

XX Sequence 212 AA;

Query Match 100.0%; Score 793; DB 2; Length 212;  
Best Local Similarity 100.0%; Pred. No. 1.2e-84;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPKKCATVLLSFLPSDFPSVRDLDTASALYEALESPEHCSPHTALRQAIL 60  
DB 30 MDIDPKKCATVLLSFLPSDFPSVRDLDTASALYEALESPEHCSPHTALRQAIL 89  
QY 61 CWGELMTLATWGVNLEDPASRLDVSVYNTNMGKFRQLLWFHISCLTFGRTVEIYL 120  
DB 90 CWGELMTLATWGVNLEDPASRLDVSVYNTNMGKFRQLLWFHISCLTFGRTVEIYL 149  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
DB 150 SFGWIRTPPAYRPPNAPILSTLPETTV 178

## RESULT 15

AAU80921

ID AAU80921 standard; protein; 212 AA.

XX AC AAU80921;

XX DT 09-APR-2002 (first entry)

XX DE Hepatitis B virus core antigen variant (HBCAg) #26.

XX KW Vaccine; molecular scaffold; pilus; pilin; HBCAg; antigen;  
XX KW hepatitis B virus capsid protein; JUN; FOS; HIV gp140;  
XX KW measles virus N protein; bee venom phospholipase; Th type 2 T-helper;  
XX KW Th2; Sinbis virus E2 protein; amyloid beta; influenza M2 antigen;  
XX KW human immunodeficiency virus infection; viral hepatitis; measles;  
XX KW chicken pox; pneumonia; tuberculosis; syphilis; malaria; allergy; cancer;  
XX KW chronic disease; arthritis; colitis; diabetes; multiple sclerosis.

XX OS Hepatitis B virus.

XX PN WO200185208-A2.

XX PD 15-NOV-2001.

XX PF 02-MAY-2001; 2001WO-IB000741.

XX PR 05-MAY-2000; 2000US-0202341P.

XX PA (CYTO-) CYTOS BIOTECHNOLOGY AG.

XX PA (SEBB) SEBBEL P.

XX PA (DUNA) DUNANT N.

XX PA (BACH) BACHMANN M.

XX PA (TISS) TISSOT A.

XX PA (LECH) LECHENER F.

XX SEBBEL P, Dunant N, Bachmann M, Tissot A, Lechener F;

XX WPI; 2002-055561/07.

XX New composition, useful for vaccine production, comprises antigen or  
XX antigenic determinant and non-natural molecular scaffold comprising  
XX organizer and core particle such as bacterial pilus or pilin protein.

XX Disclosure; Page 227-228; 287pp; English.

XX The invention relates to a composition comprising: (a) a non-natural  
XX molecular scaffold (molecular scaffold) which comprises a core particle  
XX such as a bacterial pilus or pilin protein, a recombinant form of the  
XX protein, a virus-like particle or a hepatitis B virus capsid protein  
XX (HBCAg), and an organizer; and (b) an antigen or antigenic determinant,  
XX where the molecular scaffold and antigenic determinant interact to form  
XX an ordered and repetitive antigen array. Suitable antigenic determinants  
XX include JUN, FOS, HIV gp140, measles virus N protein, bee venom  
XX phospholipase, Sinbis virus E2 protein, amyloid beta derived peptides and  
XX influenza M2 antigen. The composition (or vaccine) is useful for  
XX immunisation, by administration to a subject, where the administration  
XX produces an immune response, such as humoral, cellular or protective  
XX immune response, preferably a Th type 2 T-helper (Th2) response that is  
XX specific for the antigenic determinant. The administration induces  
XX antibodies specific for the antigenic determinant of a subtype  
XX corresponding to the Th2 subtype in the subject. The subject does not  
XX generate a Th2 subtype that is specific for pilus or pilin polypeptide or  
XX antigenic determinant. The composition is useful for the production of  
XX vaccines for prevention of infectious diseases such as human  
XX immunodeficiency virus, viral hepatitis, measles, chicken pox, pneumonia,  
XX tuberculosis, syphilis, malaria, and for treating allergy, cancer, and  
XX chronic diseases induced or accelerated by a Th1 type immune response,  
XX such as arthritis, colitis, diabetes and multiple sclerosis. The  
XX composition is useful to generate defined self-specific antibodies and  
XX specific immune responses of the Th2 type and allows the creation of  
XX highly efficient vaccines against infectious diseases, and for treating

CC allergy, cancer, and chronic diseases induced or accelerated by a Th1  
CC type immune response. The present sequence is a peptide or protein  
CC incorporated into the compositions of the invention

XX  
SQ Sequence 212 AA;  
Query Match 100.0%; Score 793; DB 5; Length 212;  
Best Local Similarity 100.0%; Pred. No. 1.2e-84;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALESPHCHSPHTALRQAIL 60  
Db 30 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALESPHCHSPHTALRQAIL 89  
QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWPHISCLTFGRETVEYLV 120  
Db 90 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWPHISCLTFGRETVEYLV 149  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

Search completed: April 23, 2004, 16:28:16  
Job time : 58 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2004 CompuGen Ltd.

OM protein - protein search, using sw model  
Run on: April 23, 2004, 16:27:14 ; Search time 23 Seconds  
(without alignments)  
334.447 Million cell updates/sec

Title: US-09-931-325C-170\_COPY\_1\_149  
Perfect score: 733  
Sequence: 1 MDDPYKFGATVLLSFLP.....PAYRPNAPILSLPTTVV 149

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

- Database : Issued Patents AA:\*
- 1: /cgn2\_6/ptodata/2/iaa/5A\_COMB.pep:\*
  - 2: /cgn2\_6/ptodata/2/iaa/5B\_COMB.pep:\*
  - 3: /cgn2\_6/ptodata/2/iaa/6A\_COMB.pep:\*
  - 4: /cgn2\_6/ptodata/2/iaa/6B\_COMB.pep:\*
  - 5: /cgn2\_6/ptodata/2/iaa/PTUS\_COMB.pep:\*
  - 6: /cgn2\_6/ptodata/2/iaa/backfiles.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	793	100.0	159	3	US-08-445-585-3
2	793	100.0	183	3	US-09-248-588-2
3	793	100.0	194	3	US-08-968-747-18
4	793	100.0	212	3	US-08-968-747-3
5	793	100.0	212	4	US-10-104-966-11
6	793	100.0	289	5	PCT-US96-10602-8
7	793	100.0	346	1	US-08-105-483-217
8	793	100.0	346	1	US-08-709-209-217
9	793	100.0	346	1	US-08-458-101-217
10	793	100.0	397	5	PCT-US96-10602-6
11	787	99.2	183	5	PCT-US96-10602-12
12	785	99.0	351	5	PCT-US96-10602-4
13	778	98.1	183	3	US-08-968-747-20
14	778	98.1	193	3	US-08-968-747-2
15	775	97.7	185	1	US-07-739-643-2
16	775	97.7	185	1	US-07-739-643-2
17	775	97.7	185	1	US-07-739-142-2
18	775	97.7	185	4	US-09-851-120-6
19	772	97.4	185	3	US-09-248-588-6
20	771	97.2	185	1	US-07-739-642-4
21	771	97.2	185	1	US-07-739-643-4
22	771	97.2	185	1	US-07-739-142-4
23	770	97.1	154	3	US-08-968-747-1
24	770	97.1	153	3	US-08-968-747-17
25	770	97.1	161	3	US-08-968-747-19
26	767.5	96.8	199	3	US-08-968-747-21
27	767	96.7	214	1	US-07-739-642-12

28	767	96.7	214	1	US-07-739-643-12	Sequence 12, Appl
29	767	96.7	214	1	US-07-739-142-12	Sequence 12, Appl
30	763	96.2	185	1	US-07-739-642-10	Sequence 10, Appl
31	763	96.2	185	1	US-07-739-643-10	Sequence 10, Appl
32	763	96.2	185	1	US-07-739-142-10	Sequence 10, Appl
33	760	95.8	183	3	US-09-248-588-4	Sequence 4, Appl
34	760	95.8	185	1	US-07-739-642-8	Sequence 8, Appl
35	760	95.8	185	1	US-07-739-643-8	Sequence 8, Appl
36	760	95.8	185	1	US-07-739-142-8	Sequence 6, Appl
37	757	95.5	214	1	US-07-739-642-6	Sequence 6, Appl
38	757	95.5	214	1	US-07-739-643-6	Sequence 6, Appl
39	757	95.5	214	1	US-07-739-142-6	Sequence 6, Appl
40	746	94.1	212	4	US-09-719-528A-4	Sequence 4, Appl
41	734.5	92.6	211	6	5196194-13	Patent No. 5196194
42	538	67.8	188	3	US-09-248-588-7	Sequence 7, Appl
43	534.5	67.4	217	3	US-09-248-588-9	Sequence 9, Appl
44	529	66.7	346	5	PCT-US96-10602-2	Sequence 2, Appl
45	132	16.6	305	3	US-09-248-588-11	Sequence 11, Appl

ALIGNMENTS

RESULT 1  
US-08-445-585-3  
; Sequence 3, Application US/08445585  
; Patent No. 6277631  
; GENERAL INFORMATION:  
; APPLICANT: No. 6277631h, Michael  
; APPLICANT: Broeker, Michael  
; TITLE OF INVENTION: Recombinant Proteins with the  
; TITLE OF INVENTION: Immunoreactivity of Hepatitis B Virus E Antigen (HBeAg), A  
; TITLE OF INVENTION: Process for the Preparation Thereof and the Use Thereof in  
; TITLE OF INVENTION: Immunoassays and Vaccines  
; NUMBER OF SEQUENCES: 12  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &  
; ADDRESSEE: Dunner  
; STREET: 1300 I Street, N.W., Suite 700  
; CITY: Washington  
; STATE: D.C.  
; COUNTRY: USA  
; ZIP: 20005-3315  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: Patent In Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/445,585  
; FILING DATE: 22-MAY-1995  
; CLASSIFICATION: 435  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US 07/846,194  
; FILING DATE: 06-MAR-1992  
; APPLICATION NUMBER: DE P 41 07 612.5  
; FILING DATE: 03-SEP-1991  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Ogden, Stasia L.  
; REGISTRATION NUMBER: P-36,228  
; REFERENCE/DOCKET NUMBER: 02481.1162-00000  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 202-408-4000  
; TELEFAX: 202-408-4400  
; INFORMATION FOR SEQ ID NO: 3:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 159 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-445-585-3

Query Match 100.0%; Score 793; DB 3; Length 159;

Best Local Similarity 100.0%; Pred. No. 3.1e-87; Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 11 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 70

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
DB 71 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 130

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 131 SFGWIRTPPAYRPPNAPILSTLPETTVV 159

RESULT 2  
US-09-248-588-2  
; Sequence 2, Application US/09248588  
; Patent No. 6231864  
; GENERAL INFORMATION:  
; APPLICANT: Birkett, Ashley J.  
; TITLE OF INVENTION: Strategically Modified Hepatitis B Core Proteins and  
; FILE REFERENCES: SYN-101 4564/69529  
; CURRENT APPLICATION NUMBER: US/09/248,588  
; EARLIER FILING DATE: 1999-02-11  
; EARLIER APPLICATION NUMBER: 60/074537  
; NUMBER OF SEQ ID NOS: 113  
; SOFTWARE: Patent In Ver. 2.0  
; SEQ ID NO 2  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
US-09-248-588-2

Query Match 100.0%; Score 793; DB 3; Length 183;  
Best Local Similarity 100.0%; Pred. No. 3.7e-87; Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 3  
US-08-968-747-18  
; Sequence 18, Application US/08968747  
; Patent No. 6060595  
; GENERAL INFORMATION:  
; APPLICANT: Scaglioni et al.  
; TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION  
; NUMBER OF SEQUENCES: 21  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Fish & Richardson P.C.  
; STREET: 225 Franklin Street  
; CITY: Boston  
; STATE: MA  
; COUNTRY: USA  
; ZIP: 02110-2804  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: Windows

SOFTWARE: Patent In Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
; Application Number: US/08/968,747  
; Filing Date: 03-SEP-1997  
; CLASSIFICATION: 435  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Fraser, Janis K.  
; REGISTRATION NUMBER: 34,819  
; REFERENCE/DOCKET NUMBER: 08472/705001  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 617/542-5070  
; TELEFAX: 617/542-8906  
; TELEX: 200154  
; INFORMATION FOR SEQ ID NO: 18:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 194 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-968-747-18

Query Match 100.0%; Score 793; DB 3; Length 194;  
Best Local Similarity 100.0%; Pred. No. 4.1e-87; Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 12 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 71

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
DB 72 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 131

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 132 SFGWIRTPPAYRPPNAPILSTLPETTVV 160

RESULT 4  
US-08-968-747-3  
; Sequence 3, Application US/08968747  
; Patent No. 6060595  
; GENERAL INFORMATION:  
; APPLICANT: Scaglioni et al.  
; TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION  
; NUMBER OF SEQUENCES: 21  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Fish & Richardson P.C.  
; STREET: 225 Franklin Street  
; CITY: Boston  
; STATE: MA  
; COUNTRY: USA  
; ZIP: 02110-2804  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: Windows  
; SOFTWARE: Patent In Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
; Application Number: US/08/968,747  
; Filing Date: 03-SEP-1997  
; CLASSIFICATION: 435  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Fraser, Janis K.  
; REGISTRATION NUMBER: 34,819  
; REFERENCE/DOCKET NUMBER: 08472/705001  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 617/542-5070  
; TELEFAX: 617/542-8906  
; TELEX: 200154  
; INFORMATION FOR SEQ ID NO: 3:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 212 amino acids

TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-968-747-3

Query Match 100.0%; Score 793; DB 3; Length 212;  
Best Local Similarity 100.0%; Pred. No. 4.6e-87;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSRLDLDTSALYREALSPHCHSPHHTALRQAIL 60  
DB 30 MDIDPYKEFGATVELLSFLPSDFPVSRLDLDTSALYREALSPHCHSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 150 SFGVWIRTPPAYRPPNAPILSTLPETTVV 178

RESULT 5  
US-10-104-966-11  
Sequence 11, Application US/10104966  
Patent No. 6880059  
GENERAL INFORMATION:  
APPLICANT: Matti Sallberg  
APPLICANT: Catharina Hultgren  
TITLE OF INVENTION: VACCINES CONTAINING RIBAVIRIN AND  
TITLE OF INVENTION: METHODS OF USE THEREOF  
FILE REFERENCE: TRIPEP 23AUSC1  
CURRENT APPLICATION NUMBER: US/10/104,966  
PRIOR FILING DATE: 2002-03-22  
PRIOR APPLICATION NUMBER: 09/705,547  
PRIOR FILING DATE: 2000-11-03  
PRIOR APPLICATION NUMBER: 60/229,175  
PRIOR FILING DATE: 2000-08-29  
NUMBER OF SEQ ID NOS: 15  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 11  
LENGTH: 212  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: Hepatitis B virus C antigen and e antigen  
OTHER INFORMATION: (HbcAg/HbeAg) sequence  
US-10-104-966-11

Query Match 100.0%; Score 793; DB 4; Length 212;  
Best Local Similarity 100.0%; Pred. No. 4.6e-87;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSRLDLDTSALYREALSPHCHSPHHTALRQAIL 60  
DB 30 MDIDPYKEFGATVELLSFLPSDFPVSRLDLDTSALYREALSPHCHSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 150 SFGVWIRTPPAYRPPNAPILSTLPETTVV 178

RESULT 6  
PCT-US96-10602-8  
Sequence 8, Application PC/TUS9610602  
GENERAL INFORMATION:  
APPLICANT: The General Hospital Corporation  
TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION  
NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:  
ADDRESSEE: Fish & Richardson P.C.  
STREET: 225 Franklin Street  
CITY: Boston  
STATE: MA  
COUNTRY: USA  
ZIP: 02110-2804  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: PCT/US96/10602  
FILING DATE:  
CLASSIFICATION:  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 60/017,814  
FILING DATE: 20-JUN-1995  
CLASSIFICATION:  
ATTORNEY/AGENT INFORMATION:  
NAME: Clark, Paul T.  
REGISTRATION NUMBER: 30,162  
REFERENCE/DOCKET NUMBER: 00786/282001  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 617/542-5070  
TELEFAX: 617/542-8906  
TELEX: 200154  
INFORMATION FOR SEQ ID NO: 8:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 289 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
PCT-US96-10602-8

Query Match 100.0%; Score 793; DB 5; Length 289;  
Best Local Similarity 100.0%; Pred. No. 7.1e-87;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSRLDLDTSALYREALSPHCHSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPVSRLDLDTSALYREALSPHCHSPHHTALRQAIL 60

QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 7  
US-08-105-483-217  
Sequence 217, Application US/08105483  
Patent No. 5494807  
GENERAL INFORMATION:  
APPLICANT: Faoletti, Enzo  
TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE  
TITLE OF INVENTION: STRAIN  
NUMBER OF SEQUENCES: 462  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Curtis, Morris & Safford  
STREET: c/o William S. Frommer  
CITY: New York  
STATE: NY  
COUNTRY: USA  
ZIP: 10036  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible

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/ OPERATING SYSTEM: PC-DOS/MS-DOS
/ SOFTWARE: Patent In Release #1.0, Version #1.25
/ CURRENT APPLICATION DATA:
/ APPLICATION NUMBER: US/08/105,483
/ FILING DATE: 12-AUG-1993
/ CLASSIFICATION: 424
/ PRIOR APPLICATION DATA:
/ APPLICATION NUMBER: US 07/847,951
/ FILING DATE: 06-MAR-1992
/ ATTORNEY/AGENT INFORMATION:
/ NAME: Frommer, William S.
/ REGISTRATION NUMBER: 25,506
/ REFERENCE/DOCKET NUMBER: 454310-2400
/ TELECOMMUNICATION INFORMATION:
/ TELEPHONE: (212) 840-3333
/ TELEFAX: (212) 840-0712
/ INFORMATION FOR SEQ ID NO: 217:
/ SEQUENCE CHARACTERISTICS:
/ LENGTH: 346 amino acids
/ TYPE: amino acid
/ STRANDEDNESS: single
/ TOPOLOGY: linear
/ US-08-105-483-217

Query Match 100.0%; Score 793; DB 1; Length 346;
Best Local Similarity 100.0%; Pred. No. 9,1e-87;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
DB 164 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 223
QY 61 CWGELMTLATWGVNLEDPASRDLVVSYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 120
DB 224 CWGELMTLATWGVNLEDPASRDLVVSYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 283
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149
DB 284 SFGWIRTPPAYRPPNAPILSTLPETTVV 312

RESULT 8
US-08-709-209-217
; Sequence 217, Application US/08/709209
; Patent No. 5762938
; GENERAL INFORMATION:
; APPLICANT: Paoletti, Enzo
; TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
; TITLE OF INVENTION: STRAIN
; NUMBER OF SEQUENCES: 462
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Curtis, Morris & Safford
; STREET: 530 Fifth Avenue
; CITY: New York
; STATE: NY
; COUNTRY: USA
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/709,209
; FILING DATE: 21-AUG-1996
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/105,483
; FILING DATE: 12-AUG-1993
; APPLICATION NUMBER: US 07/847,951
; FILING DATE: 06-MAR-1992
; ATTORNEY/AGENT INFORMATION:

NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2400
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 217:
SEQUENCE CHARACTERISTICS:
LENGTH: 346 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-105-483-217

Query Match 100.0%; Score 793; DB 1; Length 346;
Best Local Similarity 100.0%; Pred. No. 9,1e-87;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
DB 164 MDIDPYKEGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 223
QY 61 CWGELMTLATWGVNLEDPASRDLVVSYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 120
DB 224 CWGELMTLATWGVNLEDPASRDLVVSYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 283
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149
DB 284 SFGWIRTPPAYRPPNAPILSTLPETTVV 312

RESULT 9
US-08-458-101-217
; Sequence 217, Application US/08458101
; Patent No. 5766599
; GENERAL INFORMATION:
; APPLICANT: Paoletti, Enzo
; APPLICANT: Perkus, Marion E.
; APPLICANT: Taylor, Jill
; APPLICANT: Tartaglia, James
; APPLICANT: No. 5766599ton, Elizabeth K.
; APPLICANT: Riviere, Michel
; APPLICANT: de Taisne, Charles
; APPLICANT: Limbach, Keith J.
; APPLICANT: Johnson, Gerard E.
; APPLICANT: Pincus, Steven E.
; APPLICANT: Cox, William I.
; APPLICANT: Audonnet, Jean-Christophe Francis
; APPLICANT: Gettig, Russell Robert
; TITLE OF INVENTION: GENETICALLY ENGINEERED VACCINE
; TITLE OF INVENTION: STRAIN
; NUMBER OF SEQUENCES: 467
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Curtis, Morris & Safford
; ADDRESSEE: c/o William S. Frommer
; STREET: 530 Fifth Avenue
; CITY: New York
; STATE: NY
; COUNTRY: USA
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/458,101
; FILING DATE: 01-JUN-1995
; CLASSIFICATION: 424
; ATTORNEY/AGENT INFORMATION:
; NAME: Frommer, William S.
; REGISTRATION NUMBER: 25,506
; REFERENCE/DOCKET NUMBER: 454310-2740
```

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TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 840-3333
; TELEFAX: (212) 840-0712
; INFORMATION FOR SEQ ID NO: 217:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 346 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-458-101-217

Query Match 100.0%; Score 793; DB 1; Length 346;
Best Local Similarity 100.0%; Pred. No. 9.1e-87;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 164 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 223

QY 61 CWGELMTLATWGVNLEDPASRDLVVSVYNTNMGKFRQLLWFWHISCLTFGRTVIEYLV 120
Db 224 CWGELMTLATWGVNLEDPASRDLVVSVYNTNMGKFRQLLWFWHISCLTFGRTVIEYLV 283

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149
Db 284 SFGWIRTPPAYRPPNAPILSTLPETTVV 312

RESULT 10
PCT-US96-10602-6
; Sequence 6, Application PC/TUS9610602
; GENERAL INFORMATION:
; APPLICANT: The General Hospital Corporation
; TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 225 Franklin Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US96/10602
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/017,814
; FILING DATE: 20-JUN-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Clark, Paul T.
; REGISTRATION NUMBER: 30,162
; REFERENCE/DOCKET NUMBER: 00786/282001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617/542-5070
; TELEFAX: 617/542-5906
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 397 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
PCT-US96-10602-6

Query Match 100.0%; Score 793; DB 5; Length 397;
Best Local Similarity 100.0%; Pred. No. 1.1e-86;
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Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60

QY 61 CWGELMTLATWGVNLEDPASRDLVVSVYNTNMGKFRQLLWFWHISCLTFGRTVIEYLV 120
Db 61 CWGELMTLATWGVNLEDPASRDLVVSVYNTNMGKFRQLLWFWHISCLTFGRTVIEYLV 120

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149
Db 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 11
PCT-US96-10602-12
; Sequence 12, Application PC/TUS9610602
; GENERAL INFORMATION:
; APPLICANT: The General Hospital Corporation
; TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 225 Franklin Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US96/10602
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/017,814
; FILING DATE: 20-JUN-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Clark, Paul T.
; REGISTRATION NUMBER: 30,162
; REFERENCE/DOCKET NUMBER: 00786/282001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617/542-5070
; TELEFAX: 617/542-5906
; TELETYPE: 200154
; INFORMATION FOR SEQ ID NO: 12:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 183 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
PCT-US96-10602-12

Query Match 99.2%; Score 787; DB 5; Length 183;
Best Local Similarity 99.3%; Pred. No. 2e-86;
Matches 148; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60

QY 61 CWGELMTLATWGVNLEDPASRDLVVSVYNTNMGKFRQLLWFWHISCLTFGRTVIEYLV 120
Db 61 CWGELMTLATWGVNLEDPASRDLVVSVYNTNMGKFRQLLWFWHISCLTFGRTVIEYLV 120

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149
Db 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149
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RESULT 12
PCT-US96-10602-4
; Sequence 4, Application PC/TUS9610602
; GENERAL INFORMATION:
; APPLICANT: The General Hospital Corporation
; TITLE OF INVENTION: INHIBITION OF HEPATITIS B REPLICATION
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 225 Franklin Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent in Release #1.0, Version #1.30
; CURRENT APPLICATION NUMBER: PCT/US96/10602
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/017,814
; FILING DATE: 20-JUN-1995
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Clark, Paul T.
; REGISTRATION NUMBER: 30,162
; REFERENCE/DOCKET NUMBER: 00786/282001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617/542-5070
; TELEFAX: 617/542-8906
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 351 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
PCT-US96-10602-4

Query Match          99.0%; Score 785; DB 5; Length 351;
Best Local Similarity 98.7%; Pred. No. 8.4e-86;
Matches 147; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 60
Db 4 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 63
Qy 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWPHISCLTFGRTVIEYLV 120
Db 64 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWPHISCLTFGRTVIEYLV 123
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 124 SFGVWIRTPPAYRPPNAPILSTLPETTVI 152

RESULT 13
US-08-968-747-20
; Sequence 20, Application US/08968747
; Patent No. 6060595
; GENERAL INFORMATION:
; APPLICANT: Scaglioni et al.
; TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 225 Franklin Street
```

```
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: Windows
; SOFTWARE: Patent in Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/968,747
; FILING DATE: 03-SEP-1997
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Fraser, Janis K.
; REGISTRATION NUMBER: 34,819
; REFERENCE/DOCKET NUMBER: 08472/705001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617/542-5070
; TELEFAX: 617/542-8906
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 20:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 183 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-968-747-20

Query Match          98.1%; Score 778; DB 3; Length 183;
Best Local Similarity 98.7%; Pred. No. 2.4e-85;
Matches 147; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 60
Db 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALROAIL 60
Qy 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWPHISCLTFGRTVIEYLV 120
Db 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWPHISCLTFGRTVIEYLV 120
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 14
US-08-968-747-2
; Sequence 2, Application US/08968747
; Patent No. 6060595
; GENERAL INFORMATION:
; APPLICANT: Scaglioni et al.
; TITLE OF INVENTION: INHIBITION OF VIRAL REPLICATION
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson P.C.
; STREET: 225 Franklin Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: Windows
; SOFTWARE: Patent in Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/968,747
; FILING DATE: 03-SEP-1997
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Fraser, Janis K.
```



Matches 145; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALROAIL 60  
Db 1 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALROAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFMHISCLTFGRETVIEYLV 120  
Db 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFMHISCLTFGRETVIEYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTIV 149  
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTIV 149

Search completed: April 23, 2004, 16:30:53  
Job time : 30 secs

REGISTRATION NUMBER: 34,819  
REFERENCE/DOCKET NUMBER: 08472/705001  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 617/542-5070  
TELEFAX: 617/542-8906  
TELEX: 200154  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 193 amino acids  
TYPE: amino acid  
STRANDEDNESS: not relevant  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-968-747-2

Query Match 98.1%; Score 778; DB 3; Length 193;  
Best Local Similarity 98.7%; Pred. No. 2.5e-85;  
Matches 147; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALROAIL 60  
Db 11 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALROAIL 70  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFMHISCLTFGRETVIEYLV 120  
Db 71 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFMHISCLTFGRETVIEYLV 130  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTIV 149  
Db 131 SFGVWIRTPPAYRPPNAPILSTLPETTIV 159

RESULT 15  
US-07-739-642-2  
Sequence 2, Application US/07739642  
Patent No. 5173427  
GENERAL INFORMATION:  
APPLICANT: Mallonee, Richard L.  
TITLE OF INVENTION: Vectors And Hosts With Increased  
TITLE OF INVENTION: Expression Of Hbcag  
NUMBER OF SEQUENCES: 24  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Richard R. Rodrick  
STREET: 1 Becton Drive  
CITY: Franklin Lakes  
STATE: New Jersey  
COUNTRY: U.S.A.  
ZIP: 07417-1880  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/07/739,642  
FILING DATE: 19910801  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: Stierwalt, Brian K.  
REGISTRATION NUMBER: 33,213  
REFERENCE/DOCKET NUMBER: P-2272  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 201-848-5317  
TELEFAX: 201-848-9228  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 185 amino acids  
TYPE: AMINO ACID  
TOPOLOGY: linear  
US-07-739-642-2

Query Match 97.7%; Score 775; DB 1; Length 185;  
Best Local Similarity 97.3%; Pred. No. 5.5e-85;

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: April 23, 2004, 16:29:45 ; Search time 42 Seconds  
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Title: US-09-931-325C-170\_COPY\_1\_149  
Perfect score: 793  
Sequence: 1 MDIDPYKEFGATVLLSFLP.....PAYRPPNAPILSTLPETTV 149

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 1133595 seqs, 276475211 residues

Total number of hits satisfying chosen parameters: 1133595

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

- Database : Published Applications AA:\*
- 1: /cgn2\_6/ptodata/2/pubpaa/US07\_PUBCOMB.pep.\*
  - 2: /cgn2\_6/ptodata/2/pubpaa/FCT\_NEW\_PUB.pep.\*
  - 3: /cgn2\_6/ptodata/2/pubpaa/US06\_NEW\_PUB.pep.\*
  - 4: /cgn2\_6/ptodata/2/pubpaa/US06\_PUBCOMB.pep.\*
  - 5: /cgn2\_6/ptodata/2/pubpaa/US07\_NEW\_PUB.pep.\*
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  - 9: /cgn2\_6/ptodata/2/pubpaa/US09A\_PUBCOMB.pep.\*
  - 10: /cgn2\_6/ptodata/2/pubpaa/US09B\_PUBCOMB.pep.\*
  - 11: /cgn2\_6/ptodata/2/pubpaa/US09C\_PUBCOMB.pep.\*
  - 12: /cgn2\_6/ptodata/2/pubpaa/US09\_NEW\_PUB.pep.\*
  - 13: /cgn2\_6/ptodata/2/pubpaa/US10A\_PUBCOMB.pep.\*
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  - 15: /cgn2\_6/ptodata/2/pubpaa/US10C\_PUBCOMB.pep.\*
  - 16: /cgn2\_6/ptodata/2/pubpaa/US10\_NEW\_PUB.pep.\*
  - 17: /cgn2\_6/ptodata/2/pubpaa/US60\_NEW\_PUB.pep.\*
  - 18: /cgn2\_6/ptodata/2/pubpaa/US60\_PUBCOMB.pep.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	793	100.0	171	10	US-09-930-915A-263
2	793	100.0	183	8	US-08-785-997-38
3	793	100.0	183	10	US-09-387-340-38
4	793	100.0	183	10	US-09-848-616-99
5	793	100.0	183	10	US-09-931-325A-170
6	793	100.0	183	10	US-09-386-591-38
7	793	100.0	183	10	US-09-930-915A-247
8	793	100.0	183	12	US-10-289-456-39
9	793	100.0	183	12	US-10-274-616-1
10	793	100.0	183	14	US-10-243-739-39
11	793	100.0	183	14	US-10-244-065-39
12	793	100.0	183	14	US-10-289-454-39
13	793	100.0	183	14	US-10-050-902-99
14	793	100.0	183	14	US-10-898-99
15	793	100.0	183	14	US-10-080-299-1

16	793	100.0	183	14	US-10-082-014-1
17	793	100.0	183	14	US-10-372-076-1
18	793	100.0	183	15	US-10-346-190-39
19	793	100.0	183	15	US-10-465-811-30
20	793	100.0	212	9	US-09-929-935-11
21	793	100.0	212	10	US-09-848-616-114
22	793	100.0	212	12	US-10-289-456-54
23	793	100.0	212	12	US-10-312-045-2
24	793	100.0	212	13	US-10-104-966-11
25	793	100.0	212	14	US-10-243-739-54
26	793	100.0	212	14	US-10-244-065-54
27	793	100.0	212	14	US-10-289-454-54
28	793	100.0	212	14	US-10-050-902-114
29	793	100.0	212	14	US-10-050-898-114
30	793	100.0	212	15	US-10-346-190-54
31	793	100.0	212	15	US-10-465-811-45
32	793	100.0	289	9	US-09-812-862-8
33	793	100.0	397	9	US-09-812-862-6
34	791	99.7	183	10	US-09-848-616-102
35	791	99.7	183	12	US-10-289-456-42
36	791	99.7	183	14	US-10-243-739-42
37	791	99.7	183	14	US-10-244-065-42
38	791	99.7	183	14	US-10-289-454-42
39	791	99.7	183	14	US-10-050-902-102
40	791	99.7	183	14	US-10-050-898-102
41	791	99.7	183	15	US-10-346-190-42
42	791	99.7	183	15	US-10-465-811-33
43	790	99.6	183	10	US-09-848-616-104
44	790	99.6	183	10	US-09-848-616-105
45	790	99.6	183	12	US-10-289-456-44

ALIGNMENTS

RESULT 1  
US-09-930-915A-263  
; Sequence 263, Application US/09930915A  
; Publication No. US20030138765A1  
; GENERAL INFORMATION:  
; APPLICANT: Birkett, Ashley J.  
; TITLE OF INVENTION: IMMUNOGENIC HEC CHIMER PARTICLES HAVING ENHANCED  
; TITLE OF INVENTION: STABILITY  
; FILE REFERENCE: 4564/83501 ICC-102.2 PCT  
; CURRENT APPLICATION NUMBER: US/09/930,915A  
; PRIOR FILING DATE: 2001-08-15  
; PRIOR APPLICATION NUMBER: 60/226,867  
; PRIOR FILING DATE: 2000-08-22  
; PRIOR APPLICATION NUMBER: 60/225,843  
; PRIOR FILING DATE: 2000-08-16  
; NUMBER OF SEQ ID NOS: 313  
; SOFTWARE: Patentin ver. 2.1  
; SEQ ID NO 263  
; LENGTH: 171  
; TYPE: PRT  
; ORGANISM: Plasmodium falciparum

Query Match	100.0%;	Score	793;	DB	10;	Length	171;
Best Local Similarity	100.0%;	Pred. No.	1.3e-80;				
Matches	149;	Conservative	0;	Mismatches	0;	Indels	0;
Gaps	0;						

Qy	1	MDIDPYKEFGATVLLSFLPSDFPPSVRDLLDTASALYREALSPHCSPPHTRQAIL	60
Db	1	MDIDPYKEFGATVLLSFLPSDFPPSVRDLLDTASALYREALSPHCSPPHTRQAIL	60
Qy	61	CWGLMTLATVGVNLEDDPASRLVSVYNTNGLKFRQLLNWHISCLITGRTVIEYLV	120
Db	61	CWGLMTLATVGVNLEDDPASRLVSVYNTNGLKFRQLLNWHISCLITGRTVIEYLV	120
Qy	121	SFGWIRTTPPAYRPPNAPILSTLPETTV	149
Db	121	SFGWIRTTPPAYRPPNAPILSTLPETTV	149

; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Welsh & Katz, Ltd.  
; STREET: 120 South Riverside Plaza, 22nd Floor  
; CITY: Chicago  
; STATE: IL  
; COUNTRY: USA  
; ZIP: 60606  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/09/387,340  
; FILING DATE:  
; CLASSIFICATION:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Gamson, Edward P.  
; REGISTRATION NUMBER: 29,381  
; REFERENCE/DOCKET NUMBER: MON-102.0 6018/69242  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (312)655-1500  
; TELEFAX: (312)655-1501  
; INFORMATION FOR SEQ ID NO: 38:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 183 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; US-09-387-340-38  
; Query Match 100.0%; Score 793; DB 10; Length 183;  
; Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
; Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKPRQLLWPHFHSICLTFGRETIVIELV 120  
DB 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKPRQLLWPHFHSICLTFGRETIVIELV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPTTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPTTVV 149  
RESULT 4  
US-09-848-616-99  
; Sequence 99, Application US/09848616  
; Publication No. US20030054010A1  
; GENERAL INFORMATION:  
; APPLICANT: Sebbel, Peter  
; APPLICANT: Dunant, Nicolas  
; APPLICANT: Bachmann, Martin  
; APPLICANT: Tissot, Alain  
; APPLICANT: Lechner, Franziska  
; TITLE OF INVENTION: Molecular Antigen Array  
; FILE REFERENCE: 1700.0180002  
; CURRENT APPLICATION NUMBER: US/09/848,616  
; CURRENT FILING DATE: 2001-03-05  
; NUMBER OF SEQ ID NOS: 186  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 99  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
; US-09-848-616-99  
Query Match 100.0%; Score 793; DB 10; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

RESULT 2  
US-08-785-997-38  
; Sequence 38, Application US/08785997  
; Publication No. US20030021804A1  
; GENERAL INFORMATION:  
; APPLICANT: Needleman, Philip  
; TITLE OF INVENTION: An Immunological Process for Increasing  
; TITLE OF INVENTION: the HDL Cholesterol Concentration  
; NUMBER OF SEQUENCES: 50  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Welsh & Katz, Ltd.  
; STREET: 120 South Riverside Plaza, 22nd Floor  
; CITY: Chicago  
; STATE: IL  
; COUNTRY: USA  
; ZIP: 60606  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/785,997  
; FILING DATE:  
; CLASSIFICATION: 424  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Gamson Ph.D., Edward P.  
; REGISTRATION NUMBER: 29,381  
; REFERENCE/DOCKET NUMBER: MON-101.0 6018/69346  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (312)655-1500  
; TELEFAX: (312)655-1501  
; INFORMATION FOR SEQ ID NO: 38:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 183 amino acids  
; TYPE: amino acid  
; STRANDEDNESS: single  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
; US-08-785-997-38  
; Query Match 100.0%; Score 793; DB 8; Length 183;  
; Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
; Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKPRQLLWPHFHSICLTFGRETIVIELV 120  
DB 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKPRQLLWPHFHSICLTFGRETIVIELV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPTTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPTTVV 149  
RESULT 3  
US-09-387-340-38  
; Sequence 38, Application US/09387340  
; Publication No. US20030026808A1  
; GENERAL INFORMATION:  
; APPLICANT: Needleman, Philip  
; APPLICANT: Glenn, Kevin  
; APPLICANT: Krul, Elaine  
; APPLICANT: Gamson, Edward P.  
; TITLE OF INVENTION: An Immunological Process and Constructs  
; TITLE OF INVENTION: for Increasing the HDL Cholesterol Concentration  
; NUMBER OF SEQUENCES: 50

Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFTGRETVEIYLV 120  
 Db 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFTGRETVEIYLV 120  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
 Db 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 5

US-09-331-325A-170  
 ; Sequence 170, Application US/09931325A  
 ; Publication No. US20030054337A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Birkett, Ashley J.  
 ; TITLE OF INVENTION: VALERIA IMMUNOGEN AND VACCINE  
 ; FILE REFERENCE: 4564/83503 ICC-103.1  
 ; CURRENT APPLICATION NUMBER: US/09/931,325A  
 ; CURRENT FILING DATE: 2002-02-22  
 ; PRIOR APPLICATION NUMBER: 60/225,843  
 ; PRIOR FILING DATE: 2000-08-16  
 ; PRIOR APPLICATION NUMBER: USN NOT YET ASSIGNED  
 ; PRIOR FILING DATE: 2001-08-15  
 ; NUMBER OF SEQ ID NOS: 186  
 ; SOFTWARE: Patent In Ver. 2.1  
 ; SEQ ID NO 170  
 ; LENGTH: 183  
 ; TYPE: PRT  
 ; ORGANISM: Hepatitis B virus  
 US-09-331-325A-170

Query Match 100.0%; Score 793; DB 10; Length 183;  
 Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
 Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFTGRETVEIYLV 120  
 Db 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFTGRETVEIYLV 120  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
 Db 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 6

US-09-386-591-38  
 ; Sequence 38, Application US/09386591  
 ; Publication No. US20030100520A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Needleman, Philip  
 ; APPLICANT: Glenn, Kevin  
 ; TITLE OF INVENTION: An Immunological Process and Constructs  
 ; TITLE OF INVENTION: For Increasing the HDL Cholesterol Concentration by DNA  
 ; TITLE OF INVENTION: Vaccination  
 ; NUMBER OF SEQUENCES: 52  
 ; CORRESPONDENCE ADDRESS:  
 ; ADDRESSEE: Welsh & Katz, Ltd.  
 ; STREET: 120 South Riverside Plaza, 22nd Floor  
 ; CITY: Chicago  
 ; STATE: IL  
 ; COUNTRY: USA  
 ; ZIP: 60606  
 ; COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: Patent In Release #1.0, Version #1.30  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/09/386,591  
 FILING DATE:  
 CLASSIFICATION:  
 ATTORNEY/AGENT INFORMATION:  
 NAME: Gamson Ph.D., Edward P.  
 REGISTRATION NUMBER: 29,381  
 REFERENCE/DOCKET NUMBER: MON-103.0 6221/69666  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (312)655-1500  
 TELEFAX: (312)655-1501  
 INFORMATION FOR SEQ ID NO: 38:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 183 amino acids  
 TYPE: amino acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 US-09-386-591-38

Query Match 100.0%; Score 793; DB 10; Length 183;  
 Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
 Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFTGRETVEIYLV 120  
 Db 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFTGRETVEIYLV 120  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
 Db 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 7

US-09-930-915A-247  
 ; Sequence 247, Application US/09930915A  
 ; Publication No. US20030138769A1  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Birkett, Ashley J.  
 ; TITLE OF INVENTION: IMMUNOGENIC HBC CHIMER PARTICLES HAVING ENHANCED  
 ; TITLE OF INVENTION: STABILITY  
 ; FILE REFERENCE: 4564/83501 ICC-102.2 PCT  
 ; CURRENT APPLICATION NUMBER: US/09/930,915A  
 ; CURRENT FILING DATE: 2001-08-15  
 ; PRIOR APPLICATION NUMBER: 60/226,867  
 ; PRIOR FILING DATE: 2000-08-22  
 ; PRIOR APPLICATION NUMBER: 60/225,843  
 ; PRIOR FILING DATE: 2000-08-16  
 ; NUMBER OF SEQ ID NOS: 313  
 ; SOFTWARE: Patent In Ver. 2.1  
 ; SEQ ID NO 247  
 ; LENGTH: 183  
 ; TYPE: PRT  
 ; ORGANISM: Hepatitis B virus  
 US-09-930-915A-247

Query Match 100.0%; Score 793; DB 10; Length 183;  
 Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
 Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFTGRETVEIYLV 120

DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 8  
US-10-289-456-39  
; Sequence 39, Application US/10289456  
; Publication No. US20040033211A1  
; GENERAL INFORMATION:  
; APPLICANT: Bachmann, Martin  
; APPLICANT: Maurer, Patrick  
; APPLICANT: Spohn, Gunther  
; TITLE OF INVENTION: Antigen Arrays for Treatment of Bone Disease  
; FILE REFERENCE: 1700.0330001  
; CURRENT APPLICATION NUMBER: US/10/289,456  
; CURRENT FILING DATE: 2002-11-07  
; PRIOR APPLICATION NUMBER: PCT/IB02/00166  
; PRIOR FILING DATE: 2002-01-21  
; PRIOR APPLICATION NUMBER: US 10/050,902  
; PRIOR FILING DATE: 2002-01-18  
; PRIOR APPLICATION NUMBER: US 60/396,635  
; PRIOR FILING DATE: 2002-07-19  
; PRIOR APPLICATION NUMBER: US 60/331,045  
; PRIOR FILING DATE: 2001-11-07  
; NUMBER OF SEQ ID NOS: 170  
; SOFTWARE: Patent in version 3.2  
; SEQ ID NO 39  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
US-10-289-456-39

Query Match 100.0%; Score 793; DB 12; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPPHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPPHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 9  
US-10-274-616-1  
; Sequence 1, Application US/10274616  
; Publication No. US20030202982A1  
; GENERAL INFORMATION:  
; APPLICANT: Birkett, Ashley J.  
; TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACCINE  
; FILE REFERENCE: ICC 127.0 4564/88545  
; CURRENT APPLICATION NUMBER: US/10/274,616  
; CURRENT FILING DATE: 2002-10-21  
; PRIOR APPLICATION NUMBER: 09/930,915  
; PRIOR FILING DATE: 2001-08-15  
; PRIOR APPLICATION NUMBER: 10/080,299  
; PRIOR FILING DATE: 2002-02-21  
; NUMBER OF SEQ ID NOS: 87  
; SOFTWARE: Patent in version 3.1  
; SEQ ID NO 1  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus

US-10-274-616-1

Query Match 100.0%; Score 793; DB 12; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPPHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPPHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 10  
US-10-243-739-39  
; Sequence 39, Application US/10243739  
; Publication No. US20030091593A1  
; GENERAL INFORMATION:  
; APPLICANT: Bachmann, Martin F.  
; APPLICANT: Storni, Tazio  
; APPLICANT: Lechner, Franziska  
; TITLE OF INVENTION: In vivo Activation of Antigen Presenting Cells for Enhancement of  
; TITLE OF INVENTION: Immune Responses Induced by Virus Like Particles  
; FILE REFERENCE: 1700.0210001  
; CURRENT APPLICATION NUMBER: US/10/243,739  
; CURRENT FILING DATE: 2002-09-16  
; PRIOR APPLICATION NUMBER: 60/318,967  
; PRIOR FILING DATE: 2001-09-14  
; NUMBER OF SEQ ID NOS: 73  
; SOFTWARE: Patent in version 3.1  
; SEQ ID NO 39  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
US-10-243-739-39

Query Match 100.0%; Score 793; DB 14; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPPHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPPHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGKFRQLLWFHISCLTFGRETVEIYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 11  
US-10-244-065-39  
; Sequence 39, Application US/10244065  
; Publication No. US2003009668A1  
; GENERAL INFORMATION:  
; APPLICANT: Bachmann, Martin F.  
; APPLICANT: Storni, Tazio  
; APPLICANT: Maurer, Patrick  
; APPLICANT: Tissot, Alain  
; APPLICANT: Schwarz, Kacrin  
; APPLICANT: Meijerink, Edwin  
; APPLICANT: Lipowsky, Gerard  
; APPLICANT: Pumpens, Paul  
; APPLICANT: Cielens, Indulis

APPLICANT: Renhofa, Regina  
; TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particles  
; FILE REFERENCE: 1700.0220001  
; CURRENT APPLICATION NUMBER: US/10/244,065  
; CURRENT FILING DATE: 2002-09-16  
; PRIOR APPLICATION NUMBER: 60/374,145  
; PRIOR FILING DATE: 2002-04-22  
; PRIOR APPLICATION NUMBER: 60/318,994  
; PRIOR FILING DATE: 2001-09-14  
; NUMBER OF SEQ ID NOS: 73  
; SOFTWARE: Patent in version 3.1  
; SEQ ID NO 39  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
US-10-244-065-39

Query Match 100.0%; Score 793; DB 14; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 12  
US-10-289-454-39  
; Sequence 39, Application US/10289454  
; Publication No. US20030157479A1  
; GENERAL INFORMATION:  
; APPLICANT: Bachmann, Martin  
; APPLICANT: Jennings, Gary  
; APPLICANT: Sonderegger, Ivo  
; TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases  
; FILE REFERENCE: 1700.0360001  
; CURRENT APPLICATION NUMBER: US/10/289,454  
; CURRENT FILING DATE: 2003-02-10  
; PRIOR APPLICATION NUMBER: US 60/396,636  
; PRIOR FILING DATE: 2002-07-19  
; PRIOR APPLICATION NUMBER: PCT/IB02/00166  
; PRIOR FILING DATE: 2002-01-21  
; PRIOR APPLICATION NUMBER: US 10/050,902  
; PRIOR FILING DATE: 2002-01-18  
; PRIOR APPLICATION NUMBER: US 60/331,045  
; PRIOR FILING DATE: 2001-11-07  
; NUMBER OF SEQ ID NOS: 386  
; SOFTWARE: Patent in version 3.2  
; SEQ ID NO 39  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
US-10-289-454-39

Query Match 100.0%; Score 793; DB 14; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 14  
US-10-050-898-99  
; Sequence 99, Application US/10050898  
; Publication No. US20030175711A1  
; GENERAL INFORMATION:  
; APPLICANT: Renner, Wolfgang A.  
; APPLICANT: Bachmann, Martin  
; APPLICANT: Tissot, Alain  
; APPLICANT: Maurer, Patrick  
; APPLICANT: Lechner, Franziska  
; APPLICANT: Sebbel, Peter  
; APPLICANT: Piossek, Christine  
; APPLICANT: Ortmann, Rainer  
; APPLICANT: Luond, Rainer  
; APPLICANT: Staufenbiel, Matthias  
; APPLICANT: Frey, Peter  
; TITLE OF INVENTION: Molecular Antigen Array

APPLICANT: Renhofa, Regina  
; TITLE OF INVENTION: Packaging of Immunostimulatory Substances into Virus-like Particles  
; FILE REFERENCE: 1700.0220001  
; CURRENT APPLICATION NUMBER: US/10/244,065  
; CURRENT FILING DATE: 2002-09-16  
; PRIOR APPLICATION NUMBER: 60/374,145  
; PRIOR FILING DATE: 2002-04-22  
; PRIOR APPLICATION NUMBER: 60/318,994  
; PRIOR FILING DATE: 2001-09-14  
; NUMBER OF SEQ ID NOS: 73  
; SOFTWARE: Patent in version 3.1  
; SEQ ID NO 39  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
US-10-244-065-39

Query Match 100.0%; Score 793; DB 14; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 12  
US-10-289-454-39  
; Sequence 39, Application US/10289454  
; Publication No. US20030157479A1  
; GENERAL INFORMATION:  
; APPLICANT: Bachmann, Martin  
; APPLICANT: Jennings, Gary  
; APPLICANT: Sonderegger, Ivo  
; TITLE OF INVENTION: Antigen Arrays for Treatments of Allergic Eosinophilic Diseases  
; FILE REFERENCE: 1700.0360001  
; CURRENT APPLICATION NUMBER: US/10/289,454  
; CURRENT FILING DATE: 2003-02-10  
; PRIOR APPLICATION NUMBER: US 60/396,636  
; PRIOR FILING DATE: 2002-07-19  
; PRIOR APPLICATION NUMBER: PCT/IB02/00166  
; PRIOR FILING DATE: 2002-01-21  
; PRIOR APPLICATION NUMBER: US 10/050,902  
; PRIOR FILING DATE: 2002-01-18  
; PRIOR APPLICATION NUMBER: US 60/331,045  
; PRIOR FILING DATE: 2001-11-07  
; NUMBER OF SEQ ID NOS: 386  
; SOFTWARE: Patent in version 3.2  
; SEQ ID NO 39  
; LENGTH: 183  
; TYPE: PRT  
; ORGANISM: Hepatitis B virus  
US-10-289-454-39

Query Match 100.0%; Score 793; DB 14; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGKFRQLLWFWHISCLTFTGRETVEIYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 14  
US-10-050-898-99  
; Sequence 99, Application US/10050898  
; Publication No. US20030175711A1  
; GENERAL INFORMATION:  
; APPLICANT: Renner, Wolfgang A.  
; APPLICANT: Bachmann, Martin  
; APPLICANT: Tissot, Alain  
; APPLICANT: Maurer, Patrick  
; APPLICANT: Lechner, Franziska  
; APPLICANT: Sebbel, Peter  
; APPLICANT: Piossek, Christine  
; APPLICANT: Ortmann, Rainer  
; APPLICANT: Luond, Rainer  
; APPLICANT: Staufenbiel, Matthias  
; APPLICANT: Frey, Peter  
; TITLE OF INVENTION: Molecular Antigen Array

Search completed: April 23, 2004, 16:35:28  
Job time : 43 secs

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FILE REFERENCE: 1700.0190005
CURRENT APPLICATION NUMBER: US/10/050,898
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US 60/262,379
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 60/288,549
PRIOR FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: US 60/326,998
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/331,045
PRIOR FILING DATE: 2001-11-07
NUMBER OF SEQ ID NOS: 350
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 99
LENGTH: 183
TYPE: PRT
ORGANISM: Hepatitis B virus
US-10-050-898-99

Query Match      100.0%; Score 793; DB 14; Length 183;
Best Local Similarity 100.0%; Pred. No. 1.5e-80;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALRQAIL 60
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALRQAIL 60

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFHISCLTFGRETIVIELV 120
DB 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFHISCLTFGRETIVIELV 120

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 15
US-10-080-299-1
; Sequence 1, Application US/10080299
; Publication No. US20030175863A1
; GENERAL INFORMATION:
; APPLICANT: Birkett, Ashley J.
; TITLE OF INVENTION: INFLUENZA IMMUNOGEN AND VACCINE
; FILE REFERENCE: ICC 127.0 4564/84273
; CURRENT APPLICATION NUMBER: US/10/080,299
; CURRENT FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 09/930,915
; PRIOR FILING DATE: 2001-08-15
; NUMBER OF SEQ ID NOS: 83
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 183
; TYPE: PRT
; ORGANISM: Hepatitis B virus
US-10-080-299-1

Query Match      100.0%; Score 793; DB 14; Length 183;
Best Local Similarity 100.0%; Pred. No. 1.5e-80;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALRQAIL 60
DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDDTASALYREALSPHCSPHHTALRQAIL 60

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFHISCLTFGRETIVIELV 120
DB 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGLKFRQLLWFHISCLTFGRETIVIELV 120

QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149
DB 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149
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GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: April 23, 2004, 16:26:09 ; Search time 20 Seconds  
(without alignments)  
716.627 Million cell updates/sec

Title: US-09-931-325C-170\_COPY\_1\_149  
Perfect score: 793  
Sequence: 1 MDIDPYKEGATVLLSFLP.....PAYRPPNAPILSILPETTVV 149

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 283366 seqs, 96191526 residues

Total number of hits satisfying chosen parameters: 283366

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000  
Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database :  
1: PIR78:.\*  
2: PIR1:.\*  
3: PIR2:.\*  
4: PIR3:.\*  
5: PIR4:.\*

pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	793	100.0	212	1	NKVLAH
2	790	99.6	212	2	e antigen precursor
3	790	99.6	212	2	e antigen precursor
4	790	99.6	212	2	e antigen precursor
5	790	99.6	212	2	e antigen precursor
6	788	99.4	212	2	e antigen precursor
7	788	99.4	212	2	e antigen precursor
8	788	99.4	212	2	e antigen precursor
9	787	99.2	183	2	core antigen - hep
10	787	99.2	212	2	e antigen precursor
11	787	99.2	212	2	e antigen precursor
12	787	99.2	212	2	e antigen precursor
13	786	99.1	183	2	core antigen - hep
14	784	98.9	183	2	core antigen - hep
15	784	98.9	212	2	e antigen precursor
16	784	98.9	212	2	e antigen precursor
17	783	98.7	212	2	e antigen precursor
18	783	98.7	212	2	e antigen precursor
19	783	98.7	212	2	e antigen precursor
20	782	98.6	183	2	core antigen - hep
21	782	98.6	212	2	e antigen precursor
22	781	98.5	183	2	core antigen - hep
23	781	98.5	212	2	e antigen precursor
24	781	98.5	212	2	e antigen precursor
25	779	98.2	183	2	core antigen - hep
26	778	98.1	212	2	e antigen precursor
27	776	97.9	183	2	core antigen - hep
28	776	97.9	212	1	NKVLAH
29	775	97.7	212	2	e antigen precursor

30	775	97.7	214	2	S01405
31	774	97.6	183	1	NKVLA2
32	774	97.6	183	2	S0181
33	774	97.6	212	2	S25651
34	773	97.5	212	2	S53242
35	772	97.4	211	1	NKVLA1
36	772	97.4	214	1	NKVLA6
37	771	97.2	214	1	NKVLA3
38	770	97.1	214	2	S47409
39	769	97.0	212	2	S53257
40	768	96.8	183	2	S53189
41	768	96.8	212	2	S53279
42	767	96.7	212	2	S53229
43	766	96.6	183	2	S53232
44	766	96.6	183	2	S53137
45	766	96.6	183	2	S53184

ALIGNMENTS

RESULT 1

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e antigen precursor / core antigen - hepatitis B virus (subtype ayw4, isolate hb321 and  
N;Alternate names: HBe antigen precursor / HBe antigen; pre-C/C antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: subtype ayw4, isolate hb321; isolate patient Ferracuti'83; isolate patient Ca  
Cheri'83  
C;Date: 18-Dec-1981 #sequence revision 08-Nov-1996 #text change 16-Jul-1999  
C;Accession: S47405; S53191; S53209; S53234; S53264; S53269; S53277; A03711  
R;Plucieniczak, A.  
A;Description: Molecular cloning and sequencing of two complete genomes of polish isolat  
A;Reference number: S47404  
A;Accession: S47405  
A;Molecule type: DNA  
A;Residues: 1-212 <PLU>  
A;Cross-references: EMBL:Z35716; NID:G527435; PIDN:CAA84786.1; PID:G527437  
A;Experimental source: subtype ayw4, isolate hb321  
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53191  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85283; NID:G736088; PIDN:CAA59593.1; PID:G736090  
A;Experimental source: isolate patient Ferracuti'83  
A;Accession: S53209  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85290; NID:G736114; PIDN:CAA59609.1; PID:G736116  
A;Experimental source: isolate patient Casteg'83  
A;Accession: S53234  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85300; NID:G736150; PIDN:CAA59631.1; PID:G736152  
A;Experimental source: isolate patient Sanna'84  
A;Accession: S53264  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85313; NID:G736194; PIDN:CAA59659.1; PID:G736196  
A;Experimental source: isolate patient Lichezi'1'85  
A;Accession: S53249  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85306; NID:G736172; PIDN:CAA59644.1; PID:G736174  
A;Experimental source: isolate patient Flore'1'86  
A;Accession: S53262  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85312; NID:G736191; PIDN:CAA59657.1; PID:G736193  
A;Experimental source: isolate patient Lichezi'83



A/Accession: S53277  
 A/Molecule type: DNA  
 A/Residues: 30-212 <LAV>  
 A/Cross-references: EMBL:X85317; NID:G736211; PIDN:CAA59669.1; PID:G736214  
 A/Experimental source: patient Giordano-2'86  
 A/Note: due to a stop codon between the e antigen precursor  
 R;Galibert, F.; Mandart, E.; Fitoussi, F.; Tiollais, P.; Charnay, P.  
 Nature 281, 646-650, 1979  
 A/Title: Nucleotide sequence of the hepatitis B virus genome (subtype ayw) in E. coli.  
 A/Reference number: A93214; MUID:81012091; PMID:399327  
 A/Molecule type: DNA  
 A/Residues: 1-212 <GAL>  
 A/Cross-references: GB:J02203; NID:G329640; PIDN:AAA45489.1; PID:G329642  
 A/Experimental source: subtype ayw  
 C/Genetics:  
 A/Gene: C  
 C/Superfamily: hepatitis B virus core antigen  
 C/Keywords: alternative initiators; core protein  
 F;1-29/Domain: signal sequence #status predicted <SIG>  
 F;30-212/Product: core antigen #status predicted <CAG>  
 F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <BCP>  
 Query Match 100.0%; Score 793; DB 1; Length 212;  
 Best Local Similarity 100.0%; Pred. No. 2.4e-72; Mismatches 0; Indels 0; Gaps 0;  
 Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPKFEGATVLLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 60  
 DB 30 MDIDPKFEGATVLLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 89  
 QY 61 CWGELMTLATWGVNLEDPASDLVSVYNTNMGKFRQLLWPHISCLTFGRVTVEYLV 120  
 DB 90 CWGELMTLATWGVNLEDPASDLVSVYNTNMGKFRQLLWPHISCLTFGRVTVEYLV 149  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
 DB 150 SFGWIRTPPAYRPPNAPILSTLPETTV 178  
 RESULT 2  
 S53211  
 e antigen precursor / core antigen - hepatitis B virus (isolate patient Castag-1'85 and  
 N;Alternate names: HBC antigen; HBe antigen precursor / HBC antigen; pre-C/C antigen  
 N;Contains: core antigen; e antigen  
 C/Species: hepatitis B virus, HBV  
 C/Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
 C/Accession: S53211; S53197  
 R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
 submitted to the EMBL Data Library, March 1995  
 A/Reference number: S53112  
 A/Molecule type: DNA  
 A/Residues: 1-212 <LAI>  
 A/Cross-references: EMBL:X85291; NID:G736117; PIDN:CAA59611.1; PID:G736119  
 A/Experimental source: isolate patient Castag-1'85  
 A/Accession: S53197  
 A/Molecule type: DNA  
 A/Residues: 30-212 <LAW>  
 A/Cross-references: EMBL:X85284; NID:G736095; PIDN:CAA59596.1; PID:G736098  
 A/Experimental source: isolate patient Ferracuti-1'89  
 A/Note: due to a stop codon between the alternative initiators the e antigen precursor  
 C/Genetics:  
 A/Gene: C  
 C/Superfamily: hepatitis B virus core antigen  
 C/Keywords: alternative initiators; core protein  
 F;1-29/Domain: signal sequence #status predicted <SIG>  
 F;30-212/Product: core antigen #status predicted <CAG>  
 F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <BCP>

Query Match 99.6%; Score 790; DB 2; Length 212;  
 Best Local Similarity 99.3%; Pred. No. 4.9e-72;  
 Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
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 DB 30 MDIDPKFEGATVLLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 89  
 QY 61 CWGELMTLATWGVNLEDPASDLVSVYNTNMGKFRQLLWPHISCLTFGRVTVEYLV 120  
 DB 90 CWGELMTLATWGVNLEDPASDLVSVYNTNMGKFRQLLWPHISCLTFGRVTVEYLV 149  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
 DB 150 SFGWIRTPPAYRPPNAPILSTLPETTV 178  
 RESULT 3  
 S32204  
 e antigen precursor / core antigen - hepatitis B virus (subtype ayw, patients C1000 and  
 N;Alternate names: HBe antigen precursor / HBC antigen; pre-C/C antigen  
 N;Contains: core antigen; e antigen  
 C/Species: hepatitis B virus, HBV  
 C/Date: 20-Feb-1995 #sequence\_revision 20-Feb-1995 #text\_change 20-Oct-2000  
 C/Accession: S32204; S53207  
 R;Preisler-Adams, S.; Schlayer, M.J.; Peters, T.; Hettler, F.; Gerok, W.; Rasenack, J.  
 submitted to the EMBL Data Library, March 1993  
 A/Description: Identification and sequence analysis of hepatitis B virus DNA in immunolc  
 A/Reference number: S32202  
 A/Molecule type: DNA  
 A/Residues: 1-212 <PRE>  
 A/Cross-references: EMBL:X72702; NID:G288927; PIDN:CAA51257.1; PID:G288930  
 A/Experimental source: subtype ayw, patient C1000  
 R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
 submitted to the EMBL Data Library, March 1995  
 A/Reference number: S53112  
 A/Accession: S53207  
 A/Molecule type: DNA  
 A/Residues: 30-212 <LAI>  
 A/Cross-references: EMBL:X85289; NID:G736110; PIDN:CAA59607.1; PID:G736113  
 A/Experimental source: isolate patient Castaa-2'87  
 A/Note: due to a stop codon mutation between the alternative initiators the e antigen F  
 C/Genetics:  
 A/Gene: C  
 C/Superfamily: hepatitis B virus core antigen  
 C/Keywords: alternative initiators; core protein  
 F;1-29/Domain: signal sequence #status predicted <SIG>  
 F;30-212/Product: core antigen #status predicted <CAG>  
 F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <BCP>  
 Query Match 99.6%; Score 790; DB 2; Length 212;  
 Best Local Similarity 99.3%; Pred. No. 4.9e-72;  
 Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPKFEGATVLLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 60  
 DB 30 MDIDPKFEGATVLLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 89  
 QY 61 CWGELMTLATWGVNLEDPASDLVSVYNTNMGKFRQLLWPHISCLTFGRVTVEYLV 120  
 DB 90 CWGELMTLATWGVNLEDPASDLVSVYNTNMGKFRQLLWPHISCLTFGRVTVEYLV 149  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149  
 DB 150 SFGWIRTPPAYRPPNAPILSTLPETTV 178  
 RESULT 4  
 S20750  
 e antigen precursor / core antigen - hepatitis B virus (subtype ayw, patient CI)  
 N;Alternate names: HBe antigen precursor / HBC antigen; pre-C/C antigen

N;Contains: core antigen; e antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: subtype ayw, patient CI  
C;Date: 20-Feb-1995 #sequence\_revision 20-Feb-1995 #text\_change 26-Aug-1999  
C;Accession: S20750  
R;Lai, M.E.; Mazzoleni, A.P.; Balestrieri, A.; Porru, A.  
submitted to the EMBL Data Library, March 1995  
A;Description: Sequence analysis of HBV genomes isolated from patients with HBsAg negative  
A;Reference number: S20745  
A;Accession: S20750  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X65258; NID:G59434; PIDN:CAA46354.1; PID:G59436  
A;Experimental source: subtype ayw, patient CI  
C;Genetics:  
A;Gene: C  
C;Superfamily: hepatitis B virus core antigen  
C;Keywords: alternative initiators; core protein  
F;1-29/Domain: signal sequence #status predicted <SIG>  
F;30-212/Product: core antigen #status predicted <CAG>  
F;30-178/Product: e antigen #status predicted <EAG>  
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.6%; Score 790; DB 2; Length 212;  
Best Local Similarity 99.3%; Pred. No. 4.9e-72;  
Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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DB 30 MDIDPYKEGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDLYVSVYNTNMGKFLQLLWFIHISCLTFGRVTEYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDLYVSVYNTNMGKFLQLLWFIHISCLTFGRVTEYLV 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 150 AFGVWIRTPPAYRPPNAPILSTLPETTVV 178

RESULT 5  
S53200  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Ferracuti-3'91)  
N;Alternate names: HBe antigen precursor / HBC antigen; pre-C/C antigen  
N;Contains: core antigen; e antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: isolate patient Ferracuti-3'91  
C;Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C;Accession: S53200  
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53200  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85286; NID:G736101; PIDN:CAA59600.1; PID:G736103  
A;Experimental source: isolate patient Ferracuti-3'91  
C;Genetics:  
A;Gene: C  
C;Superfamily: hepatitis B virus core antigen  
C;Keywords: alternative initiators; core protein  
F;1-29/Domain: signal sequence #status predicted <SIG>  
F;30-212/Product: core antigen #status predicted <CAG>  
F;30-178/Product: e antigen #status predicted <EAG>  
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.6%; Score 790; DB 2; Length 212;  
Best Local Similarity 99.3%; Pred. No. 4.9e-72;  
Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 30 MDIDPYKEGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDLYVSVYNTNMGKFLQLLWFIHISCLTFGRVTEYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDLYVSVYNTNMGKFLQLLWFIHISCLTFGRVTEYLV 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 150 AFGVWIRTPPAYRPPNAPILSTLPETTVV 178

RESULT 6  
S53216  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Castag'3)  
N;Alternate names: HBe antigen precursor / HBC antigen; pre-C/C antigen  
N;Contains: core antigen; e antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: isolate patient Castag'3  
C;Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C;Accession: S53216  
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53216  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85293; NID:G736124; PIDN:CAA59616.1; PID:G736126  
A;Experimental source: isolate patient Castag'3  
C;Genetics:  
A;Gene: C  
C;Superfamily: hepatitis B virus core antigen  
C;Keywords: alternative initiators; core protein  
F;1-29/Domain: signal sequence #status predicted <SIG>  
F;30-212/Product: core antigen #status predicted <CAG>  
F;30-178/Product: e antigen #status predicted <EAG>  
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.4%; Score 788; DB 2; Length 212;  
Best Local Similarity 99.3%; Pred. No. 7.7e-72;  
Matches 148; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MDIDPYKEGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
DB 30 MDIDPYKEGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDLYVSVYNTNMGKFLQLLWFIHISCLTFGRVTEYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDLYVSVYNTNMGKFLQLLWFIHISCLTFGRVTEYLV 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 150 SFGVWIRTPPAYRPPNAPILSTLPETTVV 178

RESULT 7  
S53272  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Licheri-3'90)  
N;Alternate names: HBe antigen precursor / HBC antigen; pre-C/C antigen  
N;Contains: core antigen; e antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: isolate patient Licheri-3'90  
C;Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C;Accession: S53272  
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53272  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85315; NID:G736205; PIDN:CAA59664.1; PID:G736207  
A;Experimental source: isolate patient Licheri-3'90  
C;Genetics:  
A;Gene: C  
C;Superfamily: hepatitis B virus core antigen

C;Keywords: alternative initiators: core protein  
F;1-29/Domain: signal sequence #status predicted <SIG>  
F;30-212/Product: core antigen #status predicted <CAG>  
F;30-178/Product: e antigen #status predicted <EAG>  
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.4%; Score 788; DB 2; Length 212;  
Best Local Similarity 99.3%; Pred. No. 7,7e-72;  
Matches 148; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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DB 30 MDIDPYKEFGATVELLSFLPSDFPVSVDLLDTASALYREALSPHCHSPHHTALRQAIL 89  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 149  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 150 SFGVWIRTPPAYRPPNAPILSTLPETTVV 178

RESULT 8  
S53281  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Galistru-1'86)  
N;Alternate names: HBe antigen precursor / HBeC antigen; pre-C/C antigen  
N;Contains: core antigen; e antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: isolate patient Galistru-1'86  
C;Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C;Accession: S53281  
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53281  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85319; NID:g736218; PIDN:CAA59673.1; PID:g736220  
A;Experimental source: isolate patient Galistru-1'86  
C;Genetics:  
A;Gene: C  
C;Superfamily: hepatitis B virus core antigen  
C;Keywords: alternative initiators; core protein  
F;1-29/Domain: signal sequence #status predicted <SIG>  
F;30-212/Product: core antigen #status predicted <CAG>  
F;30-178/Product: e antigen #status predicted <EAG>  
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.4%; Score 788; DB 2; Length 212;  
Best Local Similarity 99.3%; Pred. No. 7,7e-72;  
Matches 148; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLLDTASALYREALSPHCHSPHHTALRQAIL 60  
DB 30 MDIDPYKEFGATVELLSFLPSDFPVSVDLLDTASALYREALSPHCHSPHHTALRQAIL 89  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 149  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 150 SFGVWIRTPPAYRPPNAPILSTLPETTVV 178

RESULT 9  
S53270  
core antigen - hepatitis B virus (isolate patient Licheri-2'87)  
N;Alternate names: HBeC antigen  
N;Contains: core antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: isolate patient Licheri-2'87

C;Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C;Accession: S53270  
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53270  
A;Molecule type: DNA  
A;Residues: 1-183 <LAI>  
A;Cross-references: EMBL:X85314; NID:g736201; PIDN:CAA59662.1; PID:g736204  
A;Experimental source: isolate patient Licheri-2'87  
A;Note: due to a stop codon between the alternative initiators the e antigen precursor  
C;Genetics:  
A;Gene: C  
C;Superfamily: hepatitis B virus core antigen  
C;Keywords: core protein

Query Match 99.2%; Score 787; DB 2; Length 183;  
Best Local Similarity 98.7%; Pred. No. 8,2e-72;  
Matches 147; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLLDTASALYREALSPHCHSPHHTALRQAIL 60  
DB 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLLDTASALYREALSPHCHSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 120  
DB 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149  
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 10  
S53225  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Chighine-2'86)  
N;Alternate names: HBe antigen precursor / HBeC antigen; pre-C/C antigen  
N;Contains: core antigen; e antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: isolate patient Chighine-2'86  
C;Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C;Accession: S53225  
R;Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53225  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85296; NID:g736137; PIDN:CAA59622.1; PID:g736139  
A;Experimental source: isolate patient Chighine-2'86  
C;Genetics:  
A;Gene: C  
C;Superfamily: hepatitis B virus core antigen  
C;Keywords: alternative initiators; core protein  
F;1-29/Domain: signal sequence #status predicted <SIG>  
F;30-212/Product: core antigen #status predicted <CAG>  
F;30-178/Product: e antigen #status predicted <EAG>  
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.2%; Score 787; DB 2; Length 212;  
Best Local Similarity 98.7%; Pred. No. 9,8e-72;  
Matches 147; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLLDTASALYREALSPHCHSPHHTALRQAIL 60  
DB 30 MDIDPYKEFGATVELLSFLPSDFPVSVDLLDTASALYREALSPHCHSPHHTALRQAIL 89  
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 120  
DB 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWPHISCLTFGRETVEIYLV 149  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTVV 149

Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

# RESULT 11

S53274  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Giordo'84)  
N/Alternate names: HBe antigen precursor / Hbc antigen; pre-C/C antigen  
N/Contains: core antigen; e antigen  
C/Species: hepatitis B virus, HBV  
A/Variety: isolate patient Giordo'84  
C/Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C/Accession: S53274  
R/Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A/Reference number: S53112  
A/Accession: S53274  
A/Molecule type: DNA  
A/Residues: 1-212 <LAI>  
A/Cross-references: EMBL:X85316; NID:g736208; PIDN:CAA59666.1; PID:g736210  
A/Experimental source: isolate patient Giordo'84  
C/Genetics:  
A/Gene: C  
C/Superfamily: hepatitis B virus core antigen  
C/Keywords: alternative initiators; core protein  
F/1-29/Domain: signal sequence #status predicted <SIG>  
F/30-212/Product: core antigen #status predicted <CAG>  
F/30-178/Product: e antigen #status predicted <EAG>  
F/179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.2%; Score 787; DB 2; Length 212;  
Best Local Similarity 99.3%; Pred. No. 9.8e-72;  
Matches 148; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
Db 1 MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
30 MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCHSPHHTALRQAIL 89  
Qy 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEIYLV 120  
Db 90 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEIYLV 149  
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

# RESULT 12

S53163  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Vittorina'92)  
N/Alternate names: HBe antigen precursor / Hbc antigen; pre-C/C antigen  
N/Contains: core antigen; e antigen  
C/Species: hepatitis B virus, HBV  
A/Variety: isolate patient Vittorina'92  
C/Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C/Accession: S53163  
R/Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A/Reference number: S53112  
A/Accession: S53163  
A/Molecule type: DNA  
A/Residues: 1-212 <LAI>  
A/Cross-references: EMBL:X85256; NID:g736050; PIDN:CAA59519.1; PID:g736052  
A/Experimental source: isolate patient Vittorina'92  
C/Genetics:  
A/Gene: C  
C/Superfamily: hepatitis B virus core antigen  
C/Keywords: alternative initiators; core protein  
F/1-29/Domain: signal sequence #status predicted <SIG>  
F/30-212/Product: core antigen #status predicted <CAG>  
F/30-178/Product: e antigen #status predicted <EAG>  
F/179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 99.2%; Score 787; DB 2; Length 212;

Best Local Similarity 99.3%; Pred. No. 9.8e-72;  
Matches 148; Conservative 0; Mismatches 1; Indels 0; Gaps 0;  
Qy 1 MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
Db 30 MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCHSPHHTALRQAIL 89  
Qy 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEIYLV 120  
Db 90 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEIYLV 149  
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

# RESULT 13

S53169  
core antigen - hepatitis B virus (isolate patient Muresu'89)  
N/Alternate names: Hbc antigen  
N/Contains: core antigen  
C/Species: hepatitis B virus, HBV  
A/Variety: isolate patient Muresu'89  
C/Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C/Accession: S53169  
R/Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A/Reference number: S53112  
A/Accession: S53169  
A/Molecule type: DNA  
A/Residues: 1-183 <LAI>  
A/Cross-references: EMBL:X85275; NID:g736057; PIDN:CAA59571.1; PID:g736060  
A/Experimental source: isolate patient Muresu'89  
A/Note: due to a stop codon between the alternative initiators the e antigen precursor  
C/Genetics:  
A/Gene: C  
C/Superfamily: hepatitis B virus core antigen  
C/Keywords: core protein

Query Match 99.1%; Score 786; DB 2; Length 183;  
Best Local Similarity 98.7%; Pred. No. 1e-71;  
Matches 147; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
Qy 1 MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
Db 1 MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCHSPHHTALRQAIL 60  
Qy 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEIYLV 120  
Db 61 CWGELMTLATWGVNLEDPASRDVLVSVYNTNMGLKFRQLLWFHISCLTFGRETVEIYLV 120  
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149

# RESULT 14

S53247  
core antigen - hepatitis B virus (isolate patient Flore-2'91)  
N/Alternate names: Hbc antigen  
N/Contains: core antigen  
C/Species: hepatitis B virus, HBV  
A/Variety: isolate patient Flore-2'91  
C/Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C/Accession: S53247  
R/Lai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A/Reference number: S53112  
A/Accession: S53247  
A/Molecule type: DNA  
A/Residues: 1-183 <LAI>  
A/Cross-references: EMBL:X85305; NID:g736168; PIDN:CAA59642.1; PID:g736171  
A/Experimental source: isolate patient Flore-2'91

A>Note: due to a stop codon between the alternative initiators the e antigen precursor

C:Genetics:  
A:Gene: C  
C:Superfamily: hepatitis B virus core antigen  
C:Keywords: core protein

Query Match 98.9%; Score 784; DB 2; Length 183;  
Best Local Similarity 98.7%; Pred.No. 1.6e-71; Mismatches 1; Indels 0; Gaps 0;  
Matches 147; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVELLSFLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
Db 1 MDIDPYKEFGATVELLSFLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
QY 61 CWGELMTLATWGVNLEDPASRLDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120  
Db 61 CWGELMTLATWGVNLEDPASRLDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETCVV 149

RESULT 15  
S53251  
e antigen precursor / core antigen - hepatitis B virus (isolate patient Flore'85)  
N;Alternate names: HBe antigen precursor / HBe antigen; pre-C/C antigen  
N;Contains: core antigen; e antigen  
C;Species: hepatitis B virus, HBV  
A;Variety: isolate patient Flore'85  
C;Date: 08-Jul-1995 #sequence\_revision 03-Aug-1995 #text\_change 26-Aug-1999  
C;Accession: S53251  
R;Jai, M.E.; Mazzoleni, A.P.; Porru, A.; Balestrieri, A.  
submitted to the EMBL Data Library, March 1995  
A;Reference number: S53112  
A;Accession: S53251  
A;Molecule type: DNA  
A;Residues: 1-212 <LAI>  
A;Cross-references: EMBL:X85307; NID:g736175; PIDN:CAAS9646.1; PID:g736177  
A;Experimental source: isolate patient Flore'85  
C:Genetics:  
A:Gene: C  
C:Superfamily: hepatitis B virus core antigen  
C:Keywords: alternative initiators; core protein  
F;1-29/Domain: signal sequence #status predicted <SIG>  
F;30-212/Product: core antigen #status predicted <CAG>  
F;30-178/Product: e antigen #status predicted <EAG>  
F;179-212/Domain: carboxyl-terminal propeptide #link EAG #status predicted <ECP>

Query Match 98.9%; Score 784; DB 2; Length 212;  
Best Local Similarity 98.7%; Pred.No. 2e-71; Mismatches 1; Indels 0; Gaps 0;  
Matches 147; Conservative 1; Mismatches 1; Indels 0; Gaps 0;  
QY 1 MDIDPYKEFGATVELLSFLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
Db 30 MDIDPYKEFGATVELLSFLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 89  
QY 61 CWGELMTLATWGVNLEDPASRLDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120  
Db 90 CWGDLTTLATWGVNLEDPASRLDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 149  
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149  
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

Search completed: April 23, 2004, 16:30:16  
Job time : 26 secs

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OM protein - protein search, using sw model

Run on: April 23, 2004, 16:22:09 ; Search time 18 Seconds

(without alignments)  
431.025 Million cell updates/sec

Title: US-09-931-325C-170\_COPY\_1\_149

Perfect score: 793  
Sequence: 1 MDIDPKYKFGATVLLSFLP.....PAYRPPNAPILSTLPETTVV 149

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 141681 seqs, 52070155 residues

Total number of hits satisfying chosen parameters: 141681

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : SwissProt\_42:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match %	Length	DB ID	Description
1	793	100.0	183	1	CORA_HPBVY
2	774	97.6	183	1	CORA_HPBVZ
3	772	97.4	185	1	CORA_HPBVM
4	772	97.4	211	1	CORA_HPBVA
5	771	97.2	185	1	CORA_HPBV2
6	761	96.0	183	1	CORA_HPBVU
7	761	96.0	183	1	CORA_HPBVO
8	760	95.8	183	1	CORA_HPBV4
9	756	95.3	214	1	CORA_HPBV9
10	754	95.1	183	1	CORA_HPBVL
11	751	94.7	212	1	CORA_HPBVT
12	749	94.5	195	1	CORA_HPBVF
13	538	67.8	188	1	CORA_HBV1
14	534.5	67.4	217	1	CORA_HPBGS
15	529	66.7	187	1	CORA_HVH8
16	132	16.6	305	1	CORA_HPBHE
17	112	14.1	305	1	CORA_HPBDB
18	112	14.1	305	1	CORA_HPBDC
19	112	14.1	305	1	CORA_HPBDM
20	110	13.9	305	1	CORA_HPBDU
21	85	10.7	306	1	RM45_MOUSE
22	79.5	10.0	3695	1	LM45_HUMAN
23	78	9.8	802	1	BCB2_ACERY
24	76.5	9.6	1226	1	YB41_METUA
25	75.5	9.5	3148	1	HD_FUGRU
26	75	9.5	784	1	DPQ2_AERPE
27	74	9.3	540	1	HXTD_YEAST
28	73.5	9.3	3587	1	SXF2_BACSU
29	73	9.2	1657	1	IQG1_HUMAN
30	72.5	9.1	306	1	RM45_HUMAN
31	72.5	9.1	2415	1	SPCA_MOUSE
32	72	9.1	377	1	YILO_YEAST
33	72	9.1	455	1	HSU1_XANAC

ALIGNMENTS

RESULT 1

ID	CORA_HPBVY	STANDARD;	PRT;	183 AA.
AC	P03146;			
DT	21-JUL-1986 (Rel. 01, Created)			
DT	21-JUL-1986 (Rel. 01, Last sequence update)			
DT	28-FEB-2003 (Rel. 41, Last annotation update)			
DE	Core antigen.			
GN	C			
OS	Hepatitis B virus (subtype ayw).			
OC	Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.			
OX	NCBI_TaxID=10418;			
RN	[1]			
RP	SEQUENCE FROM N.A.			
RX	MEDLINE=81012091; PubMed=3993327;			
RA	Galibert F., Mandart E., Fitousei F., Tiollais P., Charnay P.;			
RT	"Nucleotide sequence of the hepatitis B virus genome (subtype ayw)			
RT	Cloned in E. coli.;"			
RL	Nature 281:646-650(1979).			
RN	[2]			
RP	SEQUENCE FROM N.A.			
RC	STRAIN=Latvia;			
RX	MEDLINE=85204337; PubMed=3996597;			
RA	Bichko V., Fushko P., Dreilina D., Pumpen P., Gren E.;			
RT	"Subtype ayw variant of hepatitis B virus. DNA primary structure analysis.;"			
RL	FEBS Lett. 185:208-212(1985).			
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CC	-----			
DR	EMBL; V01460; CAA24706.1; ALT INIT.			
DR	EMBL; X02496; -; NOT ANNOTATED_CDS.			
DR	InterPro; IPR002006; Hepatitis core.			
DR	Pfam; PF00906; Hepatitis_core; 1.			
FT	Core protein; Repeat.			
FT	REPEAT 162 169			
FT	REPEAT 170 177			
FT	VARIANT 33 33			
FT	VARIANT 80 80			
FT	SEQUENCE 183 AA; 21116 MW; E0D9D9763F24E958 CRC64;			
SQ				

Query Match 100.0%; Score 793; DB 1; Length 183;  
Best Local Similarity 100.0%; Pred. No. 1.8e-72;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	MDIDPKYKFGATVLLSFLPDPFPPSVRLDPTASALYREALESPHCSPHHTALRQAIL	60
Db	1	MDIDPKYKFGATVLLSFLPDPFPPSVRLDPTASALYREALESPHCSPHHTALRQAIL	60
Qy	61	CWCELTATLWGVNLEDPASRLDVVYVNTNMGKFLQLMFWHISCLATFGRETVEYLV	120

O06596 mycobacteri  
Q19978 caenorhabdi  
P58050 arabidopsis  
Q9Y210 homo sapien  
P09917 homo sapien  
Q01738 phanerochaet  
P36304 kenedya ye  
P34633 caenorhabdi  
P58565 anabaena sp  
Q03287 escherichia  
Q06206 proteus mir  
Q9Zde9 rickettsia

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Db 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTGTRETVIEYL 120
QY 121 SFGWIRTPPAYRPPNAPILSTLPTTVV 149
Db 121 SFGWIRTPPAYRPPNAPILSTLPTTVV 149

RESULT 2
CORR HPBVZ STANDARD; PRT; 183 AA.
ID CORR HPBVZ STANDARD; PRT; 183 AA.
AC P03147;
DT 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DT 01-OCT-1989 (Rel. 12, Last annotation update)
DE Core antigen.
GN C.
OS Hepatitis B virus (subtype adw).
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10419;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=8102115; PubMed=39329;
RA Pasek M., Goto T., Gilbert W., Zink B., Schaller H., McKay P.,
RT "Hepatitis B virus genes and their expression in E. coli.";
RL Nature 282:575-579(1979).
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CC -----
CC EMBL; J02202; AAA45486.1; --
CC DR EMBL; A08967; CAA00816.1; --
CC DR PIR; B93217; NKVLAL.
CC DR InterPro; IPR002006; Hepatitis_core.
CC DA Pfam; PF00906; Hepatitis_core; 1.
CC KW Core protein; Repeat.
CC FT REPEAT 162 169
CC FT REPEAT 170 177
CC SQ SEQUENCE 183 AA; 21042 MW; 545ED0E5527F26C CRC64;

Query Match 97.6%; Score 774; DB 1; Length 183;
Best Local Similarity 96.0%; Pred. No. 1.5e-70;
Matches 143; Conservative 5; Mismatches 1; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTAAALYREALSPHCSPHHTALRQAIL 60
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTGTRETVIEYL 120
Db 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTGTRETVIEYL 120

RESULT 3
CORR HPBVW STANDARD; PRT; 185 AA.
ID CORR HPBVW STANDARD; PRT; 185 AA.
AC P03149;
DT 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DT 01-OCT-1989 (Rel. 12, Last annotation update)
DE Core antigen.
GN C.
OS Hepatitis B virus (subtype adw).
```

```
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=106821;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=83168919; PubMed=6300776;
RA Ono Y., Onda H., Sasada R., Igarashi K., Sugino Y., Nishioka K.;
RT "The complete nucleotide sequences of the cloned hepatitis B virus
RT DNA; subtype adr and adw.";
RL Nucleic Acids Res. 11:1747-1757(1983).
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CC -----
CC EMBL; V00866; --; NOT_ANNOTATED_CDS.
CC DR InterPro; IPR002006; Hepatitis_core.
CC DR Pfam; PF00906; Hepatitis_core; 1.
CC KW Core protein; Repeat.
CC FT REPEAT 164 171
CC FT REPEAT 172 179
CC SQ SEQUENCE 185 AA; 21394 MW; B86A90D541BA70F9 CRC64;

Query Match 97.4%; Score 772; DB 1; Length 185;
Best Local Similarity 96.6%; Pred. No. 2.3e-70;
Matches 144; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTGTRETVIEYL 120
Db 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWFHISCLTGTRETVIEYL 120

RESULT 4
CORR HPBVA STANDARD; PRT; 211 AA.
ID CORR HPBVA STANDARD; PRT; 211 AA.
AC P24023;
DT 01-MAR-1992 (Rel. 21, Created)
DT 01-MAR-1992 (Rel. 21, Last sequence update)
DT 16-OCT-2001 (Rel. 40, Last annotation update)
DE Core antigen.
GN C.
OS Hepatitis B virus (strain alphas).
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10411;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=90266476; PubMed=2345966;
RA Tong S., Li J., Vitvitski L., Trepo C.;
RT "Active hepatitis B virus replication in the presence of anti-HBe is
RT associated with viral variants containing an inactive pre-C region.";
RL Virology 176:596-603(1990).
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CC -----
CC EMBL; M32138; --; NOT_ANNOTATED_CDS.
CC DR PIR; A34773; NKVLAL.
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DR InterPro: IPR002006; Hepatitis core.
DR Pfam: PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 183 190
FT REPEAT 198 206
SQ SEQUENCE 211 AA; 24208 MW; B774AC72E65C75AB CRC64;

Query Match 97.4%; Score 772; DB 1; Length 211;
Best Local Similarity 96.6%; Pred. No. 2.7e-70;
Matches 144; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 29 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 88
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120
Db 89 CWGLITLTWVGNNLEDPTRDLVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 148
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 149 SFGVWIRTPPAYRPPNAPILSTLPETTV 177

RESULT 5
CORA_HPBV2
ID CORA_HPBV2 STANDARD; PRT; 185 AA.
AC P03148;
DT 21-JUL-1986 (Rel. 01, Created)
DT 01-AUG-1992 (Rel. 23, Last sequence update)
DT 15-DEC-1998 (Rel. 37, Last annotation update)
DE Core antigen.
GN Hepatitis B virus (subtype adw2).
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10408;
RN [1]
RP Valenzuela P., Quiroga M., Zaldivar J., Gray P., Rutter W.J.;
RA (In) Field B.N., Jaenisch R., Fox C.F. (eds.);
RL Animal virus genetics, pp.57-70, Academic Press, New York (1980).
DR InterPro: IPR002006; Hepatitis_core.
DR Pfam: PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 164 171
FT REPEAT 172 179
SQ SEQUENCE 185 AA; 21304 MW; 31F4DC338B507E19 CRC64;

Query Match 97.2%; Score 771; DB 1; Length 185;
Best Local Similarity 96.6%; Pred. No. 2.9e-70;
Matches 144; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120
Db 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 6
CORA_HPBVJ
ID CORA_HPBVJ STANDARD; PRT; 183 AA.
AC P17391;
DT 01-AUG-1990 (Rel. 15, Created)
DT 01-AUG-1990 (Rel. 15, Last sequence update)
DT 01-FEB-1991 (Rel. 17, Last annotation update)
DE Core antigen.

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OS Hepatitis B virus (subtype adw / strain Japan/pJ2W233).
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10413;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=89010694; PubMed=3171552;
RA Okamoto H., Tsuda F., Sakugawa H., Sastroewignjo R.I., Imai M.,
RA Miyakawa Y., Mayumi M.;
RT "Typing hepatitis B virus by homology in nucleotide sequence:
RT comparison of surface antigen subtypes.";
RL J. Gen. Virol. 69:2575-2583(1988).
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CC -----
CC EMBL; D00329; -; NOT_ANNOTATED_CDS.
DR PIR; A28925; NKVLJ1.
DR InterPro: IPR002006; Hepatitis_core.
DR Pfam: PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 162 169
FT REPEAT 170 177
SQ SEQUENCE 183 AA; 21224 MW; 9FDD6B5F5AF5E160 CRC64;

Query Match 96.0%; Score 761; DB 1; Length 183;
Best Local Similarity 95.3%; Pred. No. 2.9e-69;
Matches 142; Conservative 3; Mismatches 4; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 1 MDIDPYKEFGATVLLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
QY 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120
Db 61 CWGELMTLATWGVNLEDPASRDVSVYNTNMGLKFRQLLWPHISCLTFGRETVEIYLV 120
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 7
CORA_HPEVO
ID CORA_HPEVO STANDARD; PRT; 183 AA.
AC P17392;
DT 01-AUG-1990 (Rel. 15, Created)
DT 01-AUG-1990 (Rel. 15, Last sequence update)
DT 01-FEB-1991 (Rel. 17, Last annotation update)
DE Core antigen.
OS Hepatitis B virus (subtype adw / strain Okinawa/pODW282).
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10415;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=89010694; PubMed=3171552;
RA Okamoto H., Tsuda F., Sakugawa H., Sastroewignjo R.I., Imai M.,
RA Miyakawa Y., Mayumi M.;
RT "Typing hepatitis B virus by homology in nucleotide sequence:
RT comparison of surface antigen subtypes.";
RL J. Gen. Virol. 69:2575-2583(1988).
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CC -----
DR EMBL; D00330; -, NOT ANNOTATED_CDS.
DR PIR; B28925; NKVLJ2.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 162 169
FT REPEAT 170 177
SQ SEQUENCE 183 AA; 668DB2633122930C CRC64;

Query Match          96.0%; Score 761; DB 1; Length 183;
Best Local Similarity 95.3%; Pred. No. 2.9e-69;
Matches 142; Conservative 3; Mismatches 4; Indels 0; Gaps 0;

QY 1 MDIDPKYKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 1 MDIDPKYKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
QY 61 CWGELMTLATWGVNLEDPASRLDVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120
Db 61 CWGELMTLATWGVNLEDPASRLDVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149
Db 121 SFGWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 8
COR_A_HPBV4 STANDARD; PRT; 183 AA.
ID COR_A_HPBV4
AC P03150; P03151;
DT 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DT 01-FEB-1991 (Rel. 17, Last annotation update)
DE Core antigen.
DE Core protein.
GN Hepatitis B virus (subtype adr4), and
OS Hepatitis B virus (subtype adr), and
OS Hepatitis B virus (subtype adr / strain Indonesia/pIDW420).
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
CK NCBI_TaxID=10409, 106820, 10412;
RN [1]
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=ADR;
RX MEDLINE=831168919; PubMed=6300776;
RA Ono Y., Onda H., Sasada R., Igarashi K., Sugino Y., Nishioka K.;
RT "The complete nucleotide sequences of the cloned hepatitis B virus
RT DNA; subtype adr and adr.";
RL Nucleic Acids Res. 11:1747-1757 (1983).
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=ADR4;
RX MEDLINE=83246570; PubMed=6306594;
RA Fujiyama A., Miyazono H., Kozaki C., Yoneyama T., Ohtomo N.,
RA Matsubara K.;
RT "Cloning and structural analyses of hepatitis B virus DNAs, subtype
RT adr.";
RL Nucleic Acids Res. 11:4601-4610 (1983).
RN [3]
RP SEQUENCE FROM N.A.
RC STRAIN=ADR;
RX MEDLINE=89010694; PubMed=3171552;
RA Okamoto H., Tsuda F., Sakugawa H., Sastrosoewignjo R.I., Imai M.,
RA Miyakawa Y., Mayumi M.;
RT "Typing hepatitis B virus by homology in nucleotide sequence:
RT comparison of surface antigen subtypes.";
RL J. Gen. Virol. 69:2575-2583 (1988).
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-----
COR_A_HPBV9 STANDARD; PRT; 214 AA.
ID COR_A_HPBV9
AC P17039;
DT 01-AUG-1990 (Rel. 15, Created)
DT 01-AUG-1990 (Rel. 15, Last sequence update)
DT 01-AUG-1992 (Rel. 23, Last annotation update)
DE Core antigen.
DE Core protein.
GN Hepatitis B virus (subtype adr / strain 991).
OS Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
CK NCBI_TaxID=10410;
RN [1]
RN [2]
RP SEQUENCE FROM N.A.
RC Koechel H.G., Schueler A., Lottmann S., Thomssen R.;
RL Submitted (FEB-1990) to the EMBL/GenBank/DBJ databases.
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-----
EMBL; X51970; CAA36232.1; -
PIR; S10381; NKVLKS.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 193 200
FT REPEAT 201 208
SQ SEQUENCE 214 AA; 24722 MW; 2D668333EC5AFB8C CRC64;

Query Match          95.3%; Score 756; DB 1; Length 214;
Best Local Similarity 96.0%; Pred. No. 1.1e-66;
Matches 143; Conservative 2; Mismatches 4; Indels 0; Gaps 0;

QY 1 MDIDPKYKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
Db 30 MDIDPKYKFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 89
QY 61 CWGELMTLATWGVNLEDPASRLDVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120
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Db 90 CWGELMTLATVWGNLDPASRDVNVYNTNMGKIRQLLWFRISVLTFGRETVLEYLV 149
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

RESULT 10
CORA_HPBVL STANDARD; PRT; 183 AA.
AC P12901;
DT 01-OCT-1989 (Rel. 12, Created)
DT 01-OCT-1989 (Rel. 12, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Core antigen.
GN C.
OS Hepatitis B virus (strain lsh / chimpanzee isolate).
OC Viruses; Retroviral viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10414;
RN SEQUENCE FROM N.A.
RP MEDLINE=98258473; PubMed=2838576;
RA Vaudin M., Wolstenholme A.J., Tsiquaye K.N., Zuckerman A.J.,
RA Harrison T.J.;
RT "The complete nucleotide sequence of the genome of a hepatitis B
RT virus isolated from a naturally infected chimpanzee."
RL J. Gen. Virol. 69:1383-1389(1988).
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CC -----
DR EMBL; D00220; BAA00157.1; -
DR PIR; A28885; NKVLCF.
DR PDB; 1HHH; 31-OCT-93.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat; 3D-structure.
FT REPEAT 162 169
FT REPEAT 170 177
SQ SEQUENCE 183 AA; 20999 MW; 923DCB94A33FC0B8 CRC64;

Query Match 95.1%; Score 754; DB 1; Length 183;
Best Local Similarity 94.6%; Pred. No. 1.5e-68;
Matches 141; Conservative 5; Mismatches 3; Indels 0; Gaps 0;

Qy 1 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60
Db 1 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60
Qy 61 CWGELMTLATVWGNLDPASRDVNVYNTNMGKIRQLLWFRISVLTFGRETVLEYLV 120
Db 61 CWGELMTLATVWGNLDPASRDVNVYNTNMGKIRQLLWFRISVLTFGRETVLEYLV 120
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 11
CORA_HPBVT STANDARD; PRT; 212 AA.
AC Q05495;
DT 01-NOV-1995 (Rel. 32, Created)
DT 01-NOV-1995 (Rel. 32, Last sequence update)
DT 01-NOV-1995 (Rel. 32, Last annotation update)
DE Core antigen.
GN C.
OS Hepatitis B virus (subtype adw2 variant sf).
OC Viruses; Retroviral viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=31515;
RN SEQUENCE FROM N.A.
RP MEDLINE=90169850; PubMed=2307406;
RA Bhat R.A., Ulrich P.P., Vyas G.N.;
RT "Molecular characterization of a new variant of hepatitis B virus in
RT a persistently infected homosexual man."
RL Hepatology 11:271-276(1990).
DR PIR; A37182; NKVLH3.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 174 181
```

```
OS Hepatitis B virus (subtype adw4 / strain Brazil / isolate w4B).
OC Viruses; Retroviral viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=45410;
RN SEQUENCE FROM N.A.
RP MEDLINE=93346970; PubMed=8345355;
RA Naumann H., Schaefer S., Yoshida C.F.T., Gaspar A.M.C., Repp R.,
RA Gerlich W.H.;
RT "Identification of a new hepatitis B virus (HBV) genotype from Brazil
RT that expresses HBV surface antigen subtype adw4."
RL J. Gen. Virol. 74:1627-1632(1993).
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CC -----
DR EMBL; X69798; CAA49452.1; -
DR PIR; JQ2227; JQ2227.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT DOMAIN 178 204 ARG-RICH. 3 X 5 AA REPEATS OF S-P-R-R-R.
FT REPEAT 184 203
FT REPEAT 184 188 1.
FT REPEAT 191 195 2.
FT REPEAT 199 203 3.
SQ SEQUENCE 212 AA; 24234 MW; F832610DB7C36FD2 CRC64;

Query Match 94.7%; Score 751; DB 1; Length 212;
Best Local Similarity 93.3%; Pred. No. 3.5e-68;
Matches 139; Conservative 8; Mismatches 2; Indels 0; Gaps 0;

Qy 1 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 60
Db 30 MDIDPYKEFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCHSPHHTALRQAIL 89
Qy 61 CWGELMTLATVWGNLDPASRDVNVYNTNMGKIRQLLWFRISVLTFGRETVLEYLV 120
Db 90 CWGELMTLATVWGNLDPASRDVNVYNTNMGKIRQLLWFRISVLTFGRETVLEYLV 149
Qy 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

RESULT 12
CORA_HPBVF STANDARD; PRT; 195 AA.
AC P29178;
DT 01-DEC-1992 (Rel. 24, Created)
DT 01-DEC-1992 (Rel. 24, Last sequence update)
DT 01-DEC-1992 (Rel. 24, Last annotation update)
DE Core antigen.
GN C.
OS Hepatitis B virus (subtype adw2 variant sf).
OC Viruses; Retroviral viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=31515;
RN SEQUENCE FROM N.A.
RP MEDLINE=90169850; PubMed=2307406;
RA Bhat R.A., Ulrich P.P., Vyas G.N.;
RT "Molecular characterization of a new variant of hepatitis B virus in
RT a persistently infected homosexual man."
RL Hepatology 11:271-276(1990).
DR PIR; A37182; NKVLH3.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 174 181
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FT REPEAT 182 189
SQ SEQUENCE 195 AA; 22461 MW; E2B166F879CB7C87 CRC64;

Query Match 94.5%; Score 749; DB 1; Length 195;
Best Local Similarity 94.0%; Pred. No. 5e-68;
Matches 140; Conservative 5; Mismatches 4; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVLLSPLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 13 LDIDPYKEFGATVLLSPLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 72
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 61 CWGELMTLATVGNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFGRETVEIYLV 120
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 73 CWVELMTLATVGNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFGRETVEIYLV 132
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 121 SFGWIRTPPAYRPNAPILSTLPETTV 149
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 133 SFGWIRTPPAYRPNAPILSTLPETTV 161
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:

RESULT 13
COR_A WHV1 STANDARD; PRT; 188 AA.
ID COR_A WHV1
AC P03152;
DT 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DT 15-DEC-1998 (Rel. 37, Last annotation update)
DE Core antigen.
GN C.
OS Woodchuck hepatitis virus 1 (WHV 1),
OS Woodchuck hepatitis virus 7 (WHV 7),
OS Woodchuck hepatitis virus 59 (WHV 59), and
OS Woodchuck hepatitis virus 8 (infectious clone) (WHV 8).
OC Viruses; Retroviridae; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10430, 10432, 10431, 10434;
RN [1]
RP SEQUENCE FROM N.A.
RC SPECIES=Woodchuck hepatitis virus 1;
RX MEDLINE=82216969; PubMed=7086958;
RA Galibert F., Chen T.N., Mandart E.;
RE "Nucleotide sequence of a cloned woodchuck hepatitis virus genome:
RT comparison with the hepatitis B virus sequence.";
RL J. Virol. 41:51-65(1982).
RN [2]
RP SEQUENCE FROM N.A.
RC SPECIES=Woodchuck hepatitis virus 7, and Woodchuck hepatitis virus 59;
RX MEDLINE=88101359; PubMed=3336938;
RA Cohen J.I., Miller R.H., Rosenblum B., Denniston K., Gerin J.L.,
RA Purcell R.H.;
RT "Sequence comparison of woodchuck hepatitis virus replicative forms
RT shows conservation of the genome.";
RL Virology 162:12-20(1988).
RN [3]
RP SEQUENCE FROM N.A.
RC SPECIES=Woodchuck hepatitis virus 8 (infectious clone);
RX MEDLINE=89184524; PubMed=2928306;
RA Girones R., Cote P.J., Hornbuckle W.E., Tennant B.C., Gerin J.L.,
RA Purcell R.H., Miller R.H.;
RT "Complete nucleotide sequence of a molecular clone of woodchuck
RT hepatitis virus that is infectious in the natural host.";
RL Proc. Natl. Acad. Sci. U.S.A. 86:1846-1849(1989).
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EMBL; J02442; AAA46761.1; -
EMBL; M18752; AAA46769.1; -
EMBL; M19183; AAA46765.1; -
```

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DR EMBL; J04514; AAA46772.1; -.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 167 174
FT REPEAT 175 182
SQ SEQUENCE 188 AA; 21693 MW; 1F4454D0A7B7CE42 CRC64;

Query Match 67.8%; Score 538; DB 1; Length 188;
Best Local Similarity 64.4%; Pred. No. 7.2e-47;
Matches 96; Conservative 20; Mismatches 33; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVLLSPLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 1 MDIDPYKEFGATVLLSPLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 61 CWGELMTLATVGNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFGRETVEIYLV 120
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 61 CWDELTKLIANNSSNITSEQVTTIIVNVNDTWGLKVRQLWFIHISCLTFGRETVEIYLV 120
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 121 SFGWIRTPPAYRPNAPILSTLPETTV 149
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 121 SFGWIRTPPAYRPNAPILSTLPETTV 149
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:

RESULT 14
COR_A HPGS STANDARD; PRT; 217 AA.
ID COR_A HPGS
AC P03153;
DT 21-JUL-1986 (Rel. 01, Created)
DT 21-JUL-1986 (Rel. 01, Last sequence update)
DT 01-JAN-1990 (Rel. 13, Last annotation update)
DE Core antigen.
GN C.
OS Ground squirrel hepatitis virus (GSV).
OC Viruses; Retroviridae; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10406;
RN [1]
RP SEQUENCE FROM N.A.
RX MEDLINE=84267998; PubMed=6086950;
RA Seeger C., Ganem D., Varmus H.E.;
RE "Nucleotide sequence of an infectious molecularly cloned genome of
RT ground squirrel hepatitis virus.";
RL J. Virol. 51:367-375(1984).
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CC -----
EMBL; K02715; AAA46755.1; -.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
KW Core protein; Repeat.
FT REPEAT 196 203
FT REPEAT 204 211
SQ SEQUENCE 217 AA; 25189 MW; DF489467355EC11A CRC64;

Query Match 67.4%; Score 534.5; DB 1; Length 217;
Best Local Similarity 65.1%; Pred. No. 1.9e-46;
Matches 97; Conservative 19; Mismatches 32; Indels 1; Gaps 1;

QY 1 MDIDPYKEFGATVLLSPLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 31 MDIDPYKEFGATVLLSPLSPDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 90
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 61 CWGELMTLATVGNLEDPASRDLVVSYVNTNMGLKFRQLLWFIHISCLTFGRETVEIYLV 120
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
   :|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
```

Db 91 CWEELTRLITWMSNTTEVRR-IIUDHVNTWGLKVRQLWEHLSCLTGQHTVQEFV 149  
Qy 121 SFGWIRTPPAYRPPNAPILSTLPEHTV 149  
Db 150 SFGWIRTPPAYRPPNAPILSTLPEHTV 178

RESULT 15

CORA\_WHV8 STANDARD; PRT; 187 AA.  
AC P06433;  
DT 01-JAN-1988 (Rel. 06, Created)  
DT 01-JAN-1988 (Rel. 06, Last sequence update)  
DT 01-OCT-1989 (Rel. 12, Last annotation update)  
DE Core antigen.  
DE C  
GN Woodchuck hepatitis virus 8 (WHV 8).  
OS Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.  
OC NCBI\_TaxID=10433;  
OX [1]  
RN SEQUENCE FROM N.A.  
RX MEDLINE=86062931; PubMed=3855246;  
RA Kodama K., Ogasawara N., Yoshikawa H., Murakami S.;  
RT "Nucleotide sequence of a cloned woodchuck hepatitis virus genome:  
RT evolutionary relationship between hepadnaviruses.";  
RL J. Virol. 56:978-986(1985).  
CC -----  
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CC -----  
CC EMBL: M11082; AAA19185.1; -  
DR PIR: A03714; NKVLC2.  
DR InterPro: IPR002006; Hepatitis core.  
DR Pfam: PF00906; Hepatitis\_core; 1.  
DR Core protein; Repeat.  
FT REPEAT 166 173  
FT REPEAT 174 181  
SQ SEQUENCE 187 AA; 21579 MW; D4BC446FF7163165 CRC64;

Query Match 66.7%; Score 529; DB 1; Length 187;  
Best Local Similarity 63.8%; Pred.No. 5.7e-46;  
Matches 95; Conservative 20; Mismatches 34; Indels 0; Gaps 0;  
Qy 1 MDIDPYKEFGATVELLGFPLSPDFPSVRDLDTASALYREALSPHCSPHHTALQAIL 60  
Db 1 MDIDPYKEFGSSYQLNPLFLDFPDNLAVDTATLYBELTGREHCSPHHTAIRQALV 60  
Qy 61 CWGELMTLATWGVNLEDPASRDVLVSYVNTNMGLKFRQLLWPHISCLTGTRETVEYLV 120  
Db 61 CWDELTKLIAWSSNITSEQVRTIIVNHVNDTWGLKVRQSLWFLHLSCLTGTQVQEFV 120  
Qy 121 SFGWIRTPPAYRPPNAPILSTLPEHTV 149  
Db 121 SFGWIRTPPAYRPPNAPILSTLPEHTV 149

Search completed: April 23, 2004, 16:28:46  
Job time : 19 secs

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: April 23, 2004, 16:25:39 ; Search time 39 Seconds  
(without alignments)  
1205.441 Million cell updates/sec

Title: US-09-931-325C-170\_COPY\_1\_149  
Perfect score: 793  
Sequence: 1 MDIDPKYKFGATVLLSFLP.....PAYRPPNAPILSTLPETTVV 149

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1017041 seqs, 315518202 residues

Total number of hits satisfying chosen parameters: 1017041

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : SPTREMBL 25:\*

- 1: sp\_archaea:\*
- 2: sp\_bacteria:\*
- 3: sp\_fungi:\*
- 4: sp\_human:\*
- 5: sp\_invertebrate:\*
- 6: sp\_mammal:\*
- 7: sp\_nhc:\*
- 8: sp\_organelle:\*
- 9: sp\_phage:\*
- 10: sp\_plant:\*
- 11: sp\_rodent:\*
- 12: sp\_virus:\*
- 13: sp\_vertebrate:\*
- 14: sp\_unclassified:\*
- 15: sp\_virus:\*
- 16: sp\_bacteria:\*
- 17: sp\_archaea:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	793	100.0	176	12	Q9QRQ5 hepatitis b
2	793	100.0	205	12	Q9QRQ6 hepatitis b
3	793	100.0	212	12	Q91707 hepatitis b
4	793	100.0	212	12	O11884 hepatitis b
5	793	100.0	212	12	O89656 hepatitis b
6	791	99.7	164	12	O67904 hepatitis b
7	790	99.6	183	12	O8VB88 hepatitis b
8	790	99.6	183	12	O89437 hepatitis b
9	790	99.6	183	12	O80008 hepatitis b
10	790	99.6	183	12	O80512 hepatitis b
11	790	99.6	212	12	Q91705 hepatitis b
12	790	99.6	212	12	Q67876 hepatitis b
13	790	99.6	212	12	Q68020 hepatitis b
14	790	99.6	212	12	O68012 hepatitis b
15	790	99.6	212	12	O89597 hepatitis b
16	788	99.4	212	12	Q68025 hepatitis b

17	788	99.4	212	12	O68068 hepatitis b
18	788	99.4	212	12	O68075 hepatitis b
19	787	99.2	183	12	O68066 hepatitis b
20	787	99.2	212	12	O68070 hepatitis b
21	787	99.2	212	12	O67984 hepatitis b
22	787	99.2	212	12	O91702 hepatitis b
23	787	99.2	212	12	O68032 hepatitis b
24	787	99.2	212	12	O80SD7 hepatitis b
25	786	99.1	183	12	O67989 hepatitis b
26	786	99.1	212	12	Q91704 hepatitis b
27	785	99.0	183	12	O89531 hepatitis b
28	785	99.0	212	12	O91SV7 hepatitis b
29	785	99.0	212	12	O91SV6 hepatitis b
30	785	99.0	212	12	O91SV5 hepatitis b
31	785	99.0	212	12	O91SV4 hepatitis b
32	785	99.0	212	12	O91SV3 hepatitis b
33	785	99.0	212	12	O91SV2 hepatitis b
34	785	99.0	212	12	O91SV1 hepatitis b
35	785	99.0	212	12	O91SV0 hepatitis b
36	785	99.0	212	12	O91SV0 hepatitis b
37	785	99.0	212	12	O91SV0 hepatitis b
38	785	99.0	212	12	O91SV0 hepatitis b
39	784	98.9	183	12	O91SV0 hepatitis b
40	784	98.9	183	12	O8VB86 hepatitis b
41	784	98.9	183	12	O68048 hepatitis b
42	784	98.9	212	12	O67872 hepatitis b
43	784	98.9	212	12	O67912 hepatitis b
44	784	98.9	212	12	O68051 hepatitis b
45	784	98.9	214	12	Q91708 hepatitis b

ALIGNMENTS

RESULT 1

Q9QRQ5 PRELIMINARY; PRT; 176 AA.  
ID AC Q9QRQ5; DT 01-MAY-2000 (TREMREL. 13, Created)  
DT 01-MAY-2000 (TREMREL. 13, Last sequence update)  
DT 01-OCT-2002 (TREMREL. 22, Last annotation update)  
DE Core protein (Fragment).  
GN C.  
OS Hepatitis B virus.  
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.  
OX NCBI\_TaxID=10407;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC STRAIN=Mut 50;  
RA Saxena A., Acharya S.K., Nayak B., Panda S.K.;  
RT "HBV mutants in non-A to E acute liver failure patients in India."  
RL Submitted (DEC-1998) to the EMBL/GenBank/DBSJ databases.  
DR EMBL; AFL15425; AAD49194.1; -  
DR InterPro; IPR002006; Hepatitis core.  
DR Pfam; PF00306; Hepatitis core; 1.  
DR NON\_TER 176  
SQ SEQUENCE 176 AA; 20297 MW; 28397ABF657D93F4 CRC64;

Query Match 100.0%; Score 793; DB 12; Length 176;  
Best Local Similarity 100.0%; Pred. No. 1.5e-80;  
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 1 MDIDPKYKFGATVLLSFLPDPFSPVRLDITASALYREALSPHCSPHTALRQAIL 60  
Db 1 MDIDPKYKFGATVLLSFLPDPFSPVRLDITASALYREALSPHCSPHTALRQAIL 60  
QY 61 CWGELMTLWTWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFLHSCILTFGRETVEYLV 120  
Db 61 CWGELMTLWTWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFLHSCILTFGRETVEYLV 120  
QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
Db 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

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RESULT 2
Q90R06 PRELIMINARY; PRT; 205 AA.
AC Q90R06;
DT 01-MAY-2000 (TrEMBLrel. 13, Created)
DT 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
DT 01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
DE Precore/core protein (Fragment).
GN C.
OS Hepatitis B virus.
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10407;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=Mut 50;
RA Saxena A., Acharya S.K., Navak B., Panda S.K.;
RT "HBV mutants in non-A to E acute liver failure patients in India.";
RL Submitted (DEC-1998) to the EMBL/GenBank/DBJ databases.
DR EMBL; AF115425; AAD49193.1; -.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
FT NON_TER 205
SQ SEQUENCE 205 AA; 23531 MW; ASCF2117AC9938E6 CRC64;

Query Match 100.0%; Score 793; DB 12; Length 205;
Best Local Similarity 100.0%; Pred. No. 1.9e-80;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 60
Db 30 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 120
Db 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

RESULT 3
Q91J77 PRELIMINARY; PRT; 212 AA.
AC Q91J77;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Pre-core protein.
OS Hepatitis B virus.
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10407;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=119-Fam E;
RA Zampino R., Lobello S., Chiaramonte M., Venturi-Pasini C., Dumpis U.,
RT Thursz M., Karayiannis P.;
RT "Intra-familial transmission of Hepatitis B virus in Italy;
RT Phylogenetic sequence analysis and amino acid variation of the core
RT Gene.";
RL J. Hepatol. 0:0-0(2002).
DR EMBL; AF419515; AAL15943.1; -.
DR PIR; S33686; S33686.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
FT CHAIN 30 212 CORE PROTEIN.
SQ SEQUENCE 212 AA; 24377 MW; B9C16A527905AF8 CRC64;

Query Match 100.0%; Score 793; DB 12; Length 212;
Best Local Similarity 100.0%; Pred. No. 1.9e-80;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 60
Db 30 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 120
Db 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

RESULT 5
Q89656 PRELIMINARY; PRT; 212 AA.
AC Q89656;
DT 01-NOV-1996 (TrEMBLrel. 01, Created)
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Core antigen (Precore protein) (Core and e antigen).
GN PREC OR C.

```

```

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 60
Db 30 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 120
Db 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

RESULT 4
O11884 PRELIMINARY; PRT; 212 AA.
ID O11884;
AC O11884;
DT 01-JUL-1997 (TrEMBLrel. 04, Created)
DT 01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Core antigen (Core and e antigens).
GN C.
OS Hepatitis B virus.
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10407;
RN [1]
RP SEQUENCE FROM N.A.
RA Rao B.S., Casey J.L., Rinaudo J.S., Korba B.E.;
RT "Complete nucleotide sequence of a molecular clone of hepatitis B
RT virus from the 2.2.15 cell line.";
RL Submitted (MAR-1997) to the EMBL/GenBank/DBJ databases.
RN [2]
RP SEQUENCE FROM N.A.
RC STRAIN=CH2-OHF;
RA Duong T.N., Michitaka K., Chen Y., Horike N., Kawai K., Onji M.;
RT "A comparison of genotype C and D hepatitis B virus: a clinical and
RT molecular biological study.";
RL Submitted (AUG-2002) to the EMBL/GenBank/DBJ databases.
DR EMBL; U95551; AAB59971.1; -.
DR EMBL; AB090269; BACS7440.1; -.
DR PIR; C94409; NKVLA3.
DR PIR; S33686; S33686.
DR InterPro; IPR002006; Hepatitis_core.
DR Pfam; PF00906; Hepatitis_core; 1.
SQ SEQUENCE 212 AA; 24360 MW; D301689E8B05A1B8 CRC64;

Query Match 100.0%; Score 793; DB 12; Length 212;
Best Local Similarity 100.0%; Pred. No. 1.9e-80;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 60
Db 30 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSEPHCSPHHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 120
Db 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFKRQLLWFWHISCLTFGRETVEIYL 149

QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
Db 150 SFGVWIRTPPAYRPPNAPILSTLPETTV 178

RESULT 5
Q89656 PRELIMINARY; PRT; 212 AA.
ID Q89656;
AC Q89656;
DT 01-NOV-1996 (TrEMBLrel. 01, Created)
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Core antigen (Precore protein) (Core and e antigen).
GN PREC OR C.

```

QY	1	MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCSPHHTALRQAIL	60
Db	30	MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCSPHHTALRQAIL	89
QY	61	CWGLMNTLATWGVNLEDPASRDVLVSYVNTNMGKLFKRLWFIHISCLTFGRETVEYL	120
Db	90	CWGLMNTLATWGVNLEDPASRDVLVSYVNTNMGKLFKRLWFIHISCLTFGRETVEYL	149
QY	121	SGFWIRTPPAYRPPNAPILSTLPETTV	149
Db	150	SGFWIRTPPAYRPPNAPILSTLPETTV	178

RESULT 6

Q67904	PRELIMINARY;	PRT;	164 AA.
ID	Q67904		
AC	Q67904;		
DT	01-NOV-1996 (TReMBLrel. 01, Created)		
DT	01-NOV-1996 (TReMBLrel. 01, Last sequence update)		
DT	01-DEC-2001 (TReMBLrel. 19, Last annotation update)		
DE	Nucleocapsid protein (Fragment).		
GN	HCCAG.		
OS	Hepatitis B virus.		
OC	Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.		
OX	NCBI_TaxID=10407;		
RN	[1]		
RP	SEQUENCE FROM N.A.		
RC	STRAIN=case 8;		
EX	MEDLINE=95294561; PubMed=7775955;		
RA	Valliammai T., Thyagarajan S.P., Zuckerman A.J., Harrison T.J.;		
RT	"Precore and core mutations in HBV from individuals in India with chronic infection.";		
RL	J. Med. Virol. 45:321-325(1995).		
DR	EMBL; Z68142; CAA92251.1; -.		
DR	InterPro; IPR002006; Hepatitis core.		
DR	Pfam; PF00906; Hepatitis_core; 1.		
FT	NON TBR		
FT	164		
SEQ	SEQUENCE 164 AA; 18717 MW; BAE84DDE5B8DF11 CRC64;		

Query Match 99.7%; Score 791; DB 12; Length 164;  
 Best Local Similarity 99.3%; Pred. No. 2.3e-80;  
 Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0

QY	1	MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCSPHHTALRQAIL	60
Db	1	MDIDPYKEFGATVLLSFLPSDFPSPVRDLDTASALYREALSPHCSPHHTALRQAIL	60
QY	61	CWGLMNTLATWGVNLEDPASRDVLVSYVNTNMGKLFKRLWFIHISCLTFGRETVEYL	120
Db	61	CWGLMNTLATWGVNLEDPASRDVLVSYVNTNMGKLFKRLWFIHISCLTFGRETVEYL	120
QY	121	SGFWIRTPPAYRPPNAPILSTLPETTV	149
Db	121	SGFWIRTPPAYRPPNAPILSTLPETTV	149

RESULT 7

Q8VBFB	PRELIMINARY;	PRT;	183 AA.
ID	Q8VBFB		
AC	Q8VBFB;		
DT	01-MAR-2002 (TReMBLrel. 20, Created)		
DT	01-MAR-2002 (TReMBLrel. 20, Last sequence update)		
DT	01-OCT-2002 (TReMBLrel. 22, Last annotation update)		
DE	Core protein.		
OS	Hepatitis B virus.		
OC	Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.		
OX	NCBI_TaxID=10407;		
RN	[1]		
RP	SEQUENCE FROM N.A.		
RC	STRAIN=I54;		
RA	Jazayeri M., Sran N., Gish R., Basuni A.A., Cooksley G., Locarnini S.;		
RA	Carman W.F.;		

```
Query Match      100.0%; Score 793; DB 12; Length 212;
Best Local Similarity 100.0%; Pred. No. 1.9e-80;
Matches 149; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

RT "HBV core sequence: definition of genotype-specific variability and correlation with geographic origin."  
 RL Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.  
 DR EMBL; AF324140; AAL31853.1; -  
 DR InterPro; IPR002006; Hepatitis core.  
 DR Pfam; PF00906; Hepatitis core; 1.  
 SQ SEQUENCE 183 AA; 21100 MW; 62D9CA5700042559 CRC64;

Query Match 99.6%; Score 790; DB 12; Length 183;  
 Best Local Similarity 99.3%; Pred. No. 3.4e-80;  
 Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
 DB 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
 DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 8  
 Q89437 PRELIMINARY; PRT; 183 AA.  
 AC Q89437;  
 DT 01-NOV-1996 (TREMELrel. 01, Created)  
 DT 01-NOV-1996 (TREMELrel. 01, Last sequence update)  
 DT 01-OCT-2002 (TREMELrel. 22, Last annotation update)  
 DE Core protein (X, PREC and C genes) (CASTAA 2).  
 GN CORE.  
 OS Hepatitis B virus, and  
 OS Hepatitis B virus (subtype ayw).  
 OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.  
 OX NCBI\_TaxID=10407; 10418;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RQ SPECIES=Hepatitis B virus; STRAIN=I59;  
 RA Jazayeri M., Sran N., Gish R., Basuni A.A., Cooksley G., Locarnini S., Carman W.F.;  
 RA "HBV core sequence: definition of genotype-specific variability and correlation with geographic origin."  
 RT Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.  
 RN [2]  
 RP SEQUENCE FROM N.A.  
 RC SPECIES=Hepatitis B virus (subtype ayw);  
 RA Lai M.E., Mazzoleni A.P., Porru A., Balestrieri A.;  
 RL Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.  
 RN [3]  
 RP SEQUENCE FROM N.A.  
 RC SPECIES=Hepatitis B virus (subtype ayw);  
 RA Karayiannis P.;  
 RL Submitted (DEC-1995) to the EMBL/GenBank/DDBJ databases.  
 RN [4]  
 RP SEQUENCE FROM N.A.  
 RC SPECIES=Hepatitis B virus (subtype ayw);  
 RA Karayiannis P.;  
 RL Submitted (SEP-1995) to the EMBL/GenBank/DDBJ databases.  
 DR EMBL; AF324146; AAL31859.1; -  
 DR EMBL; X85289; CAA59607.1; -  
 DR EMBL; X80925; CAA56888.1; -  
 DR InterPro; IPR002006; Hepatitis core.  
 DR Pfam; PF00906; Hepatitis core; 1.  
 SQ SEQUENCE 183 AA; 21102 MW; 2B9902063F253228 CRC64;

Query Match 99.6%; Score 790; DB 12; Length 183;  
 Best Local Similarity 99.3%; Pred. No. 3.4e-80;  
 Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60

DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
 DB 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
 DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 9  
 Q88008 PRELIMINARY; PRT; 183 AA.  
 AC Q88008;  
 DT 01-NOV-1996 (TREMELrel. 01, Created)  
 DT 01-NOV-1996 (TREMELrel. 01, Last sequence update)  
 DT 01-OCT-2002 (TREMELrel. 22, Last annotation update)  
 DE Core protein.  
 GN CORE.  
 OS Hepatitis B virus.  
 OS Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.  
 OX NCBI\_TaxID=10407;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RQ SPECIES=Hepatitis B virus; STRAIN=I59;  
 RA Lai M.E., Mazzoleni A.P., Porru A., Balestrieri A.;  
 RL Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.  
 DR EMBL; X85284; CAA59596.1; -  
 DR InterPro; IPR002006; Hepatitis core.  
 DR Pfam; PF00906; Hepatitis core; 1.  
 SQ SEQUENCE 183 AA; 21102 MW; 50B9D9763F25E958 CRC64;  
 Query Match 99.6%; Score 790; DB 12; Length 183;  
 Best Local Similarity 99.3%; Pred. No. 3.4e-80;  
 Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;  
 QY 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 DB 1 MDIDPYKEFGATVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHHTALRQAIL 60  
 QY 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
 DB 61 CWGELMTLATWGVNLEDPASRDLVVSYVNTNMGKFRQLLWFIHISCLTFGRETVEIYLV 120  
 QY 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149  
 DB 121 SFGWIRTPPAYRPPNAPILSTLPETTVV 149

RESULT 10  
 Q805R2 PRELIMINARY; PRT; 183 AA.  
 AC Q805R2;  
 DT 01-JUN-2003 (TREMELrel. 24, Created)  
 DT 01-JUN-2003 (TREMELrel. 24, Last sequence update)  
 DT 01-OCT-2003 (TREMELrel. 25, Last annotation update)  
 DE Core protein.  
 OS Hepatitis B virus.  
 OS Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.  
 OX NCBI\_TaxID=10407;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RQ SPECIES=Hepatitis B virus (subtype ayw);  
 RA Chen Y., Michitaka K., Matsubara H., Yamamoto K., Horike N., Onji M.;  
 RA "Complete genome sequence of hepatitis B virus (HBV) from a patient with fulminant hepatitis without precore and core promoter mutations: comparison with HBV from a patient with acute hepatitis infected from the same infectious source."  
 RT J. Hepatol. 38:84-90 (2003).  
 RL EMBL; AB078031; BAC54833.1; -  
 DR EMBL; AB078032; BAC54836.1; -



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DR EMBL; AB078033; BACS4839.1; -.
DR InterPro; IPR002006; Hepatitis core.
DR Pfam; PF00906; Hepatitis core; 1.
SQ SEQUENCE 183 AA; 21130 MW; 1B39CDC97BDA0319 CRC64;

Query Match          99.6%; Score 790; DB 12; Length 183;
Best Local Similarity 99.3%; Pred. No. 3.4e-80;
Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 60
DB 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 60
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 120
DB 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 120
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
DB 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149

RESULT 11
Q917J5 PRELIMINARY; PRT; 212 AA.
ID Q917J5
AC Q917J5;
DT 01-DEC-2001 (TrEMBLrel. 19, Created)
DT 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Pre-core protein.
OS Hepatitis B virus.
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10407;
RN [1]
RP SEQUENCE FROM N.A.
RC STRAIN=I21-Fam E;
RA Zampino R., Lobello S., Chiaramonte M., Venturi-Pasini C., Dumpie U.,
  Thursz M., Karayiannis P.;
RT "Intra-familial transmission of Hepatitis B virus in Italy;
  Phylogenetic sequence analysis and amino acid variation of the core
  gene.";
RL J. Hepatol. 0:0-0(2002).
DR EMBL; AF419517; AAL15945.1; -.
DR PIR; S33686; S33686.
DR InterPro; IPR002006; Hepatitis core.
DR Pfam; PF00906; Hepatitis core; 1.
FT CHAIN 30 212 CORE PROTEIN.
SQ SEQUENCE 212 AA; 24412 MW; 19D58A4EBB05B243 CRC64;

Query Match          99.6%; Score 790; DB 12; Length 212;
Best Local Similarity 99.3%; Pred. No. 4.1e-80;
Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 60
DB 30 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 89
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 120
DB 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 149

RESULT 12
Q67876 PRELIMINARY; PRT; 212 AA.
ID Q67876
AC Q67876;
DT 01-NOV-1996 (TrEMBLrel. 01, Created)
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)

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DE Pre C/C ORF.
OS Hepatitis B virus.
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10407;
RN [1]
RP SEQUENCE FROM N.A.
RA Lai M.E., Mazzoleni A.P., Balestrieri A., Melis A., Portu A.;
RT "Sequence analysis of HBV genomes isolated from patients with HBsAg
  Negative chronic liver disease.";
RL Submitted (MAR-1992) to the EMBL/GenBank/DBJ databases.
DR EMBL; X65258; CAA46354.1; -.
DR PIR; C94409; NKVLA3.
DR PIR; S20750; S20750.
DR PIR; S33686; S33686.
DR InterPro; IPR002006; Hepatitis core.
DR Pfam; PF00906; Hepatitis core; 1.
SQ SEQUENCE 212 AA; 24348 MW; 6E91CC7D068EB573 CRC64;

Query Match          99.6%; Score 790; DB 12; Length 212;
Best Local Similarity 99.3%; Pred. No. 4.1e-80;
Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 60
DB 30 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 89
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 120
DB 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 149
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149
DB 150 AFGVWIRTPPAYRPPNAPILSTLPETTV 178

RESULT 13
Q68020 PRELIMINARY; PRT; 212 AA.
ID Q68020
AC Q68020;
DT 01-NOV-1996 (TrEMBLrel. 01, Created)
DT 01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
DT 01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
DE Pre-c/core protein.
OS Hepatitis B virus.
OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
OX NCBI_TaxID=10407;
RN [1]
RP SEQUENCE FROM N.A.
RA Lai M.E., Mazzoleni A.P., Portu A., Balestrieri A.;
RL Submitted (MAR-1995) to the EMBL/GenBank/DBJ databases.
DR EMBL; X85291; CAA59611.1; -.
DR PIR; C94409; NKVLA3.
DR PIR; S33686; S33686.
DR PIR; S53211; S53211.
DR InterPro; IPR002006; Hepatitis core.
DR Pfam; PF00906; Hepatitis core; 1.
SQ SEQUENCE 212 AA; 24336 MW; 63006A4EB804A1B8 CRC64;

Query Match          99.6%; Score 790; DB 12; Length 212;
Best Local Similarity 99.3%; Pred. No. 4.1e-80;
Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 60
DB 30 MDIDPYKEFGATVELLSFLPSDFPVSVDLDTASALYREALSPHCSPHHTALRQAIL 89
QY 61 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 120
DB 90 CWGELMTLATWGVNLEDPASRDLVSVYNTNMGKFRQLLWFHISCLTFGRETIVIELV 149
QY 121 SFGVWIRTPPAYRPPNAPILSTLPETTV 149

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Db 150 SFGWIRTPPAYPPNAPILSTLPETTV 178

RESULT 14

Q68012 PRELIMINARY; PRT; 212 AA.

AC Q68012

DT 01-NOV-1996 (TREMELrel. 01, Created)

DT 01-NOV-1996 (TREMELrel. 01, Last sequence update)

DT 01-OCT-2003 (TREMELrel. 25, Last annotation update)

DE Pre-C/core protein.

GN PRE-C/CORE.

OS Hepatitis B virus.

OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.

OX NCBI\_TaxID=10407;

RN [1]

RP SEQUENCE FROM N.A.

RA Lai M.E., Mazzoleni A.P., Porru A., Balestrieri A.

RL Submitted (MAR-1995) to the EMBL/GenBank/DBJ databases.

DR EMBL; X85286; CAA59600.1; -.

DR PIR; S33686; S33686.

DR PIR; S53200; S53200.

DR InterPro; IPR002006; Hepatitis core.

DR Pfam; PF00906; Hepatitis core; 1.

DR S33686; S33686.

SQ SEQUENCE 212 AA; 24250 MW; D30179D05DCD51B8 CRC64;

Query Match 99.6%; Score 790; DB 12; Length 212;

Best Local Similarity 99.3%; Pred. No. 4.1e-80;

Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 60

Db 30 MDIDPYKFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDVVSYYNTNMGLKFRQLLWFHISCLTFCGTVEYLV 120

Db 90 CWGELMTLATWGVNLEDPASRDVVSYYNTNMGLKFRQLLWFHISCLTFCGTVEYLV 149

QY 121 SFGWIRTPPAYPPNAPILSTLPETTV 149

Db 150 SFGWIRTPPAYPPNAPILSTLPETTV 178

RESULT 15

Q89597 PRELIMINARY; PRT; 212 AA.

AC Q89597

DT 01-NOV-1996 (TREMELrel. 01, Created)

DT 01-NOV-1996 (TREMELrel. 01, Last sequence update)

DT 01-OCT-2003 (TREMELrel. 25, Last annotation update)

DE HECAG (PRE-CORE protein) (PRECORE/core protein).

OS Hepatitis B virus (subtype ayw), and

OS Hepatitis B virus.

OC Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.

OX NCBI\_TaxID=10418, 10407;

RN [1]

RP SEQUENCE FROM N.A.

RC SPECIES=Hepatitis B virus (subtype ayw); STRAIN=SUB-TYPE AYW;

RX MEDLINE=94079539; PubMed=8257295;

RA Preisler-Adams S., Schlayer M.J., Peters T., Hettler F., Gerok W.,

RA Rasenack J.

RT "Sequence analysis of hepatitis B virus DNA in immunologically

RT negative infection."

RL Arch. Virol. 133:385-396(1993).

RN [2]

RP SEQUENCE FROM N.A.

RC SPECIES=Hepatitis B virus (subtype ayw); STRAIN=AYW;

RA Karayiannis P.

RL Submitted (DEC-1995) to the EMBL/GenBank/DBJ databases.

RN [3]

RP SEQUENCE FROM N.A.

RC SPECIES=Hepatitis B virus (subtype ayw); STRAIN=AYW;

RA Karayiannis P.

Submitted (SEP-1995) to the EMBL/GenBank/DBJ databases.

(4)

RL SEQUENCE FROM N.A.

RP SPECIES=Hepatitis B virus; STRAIN=I43-Fam G;

RC Zampino R., Lobello S., Chiaramonte M., Venturi-Pasini C., Dumpis U.,

RA Thurez M., Karayiannis P.

RT "Intra-familial transmission of Hepatitis B virus in Italy:

RT Phylogenetic sequence analysis and amino acid variation of the core

RT gene."

RL J. Hepatol. 0:0-0(2002).

RN [5]

RP SEQUENCE FROM N.A.

RC SPECIES=Hepatitis B virus; STRAIN=Gaml119F29, and Gaml821F217;

RA Dumpis U., Mendy M., Karayiannis P.;

RT "Prevalence of HBV core promoter/precore/core mutations in Gambian

RT chronic carriers."

RL Submitted (FEB-2001) to the EMBL/GenBank/DBJ databases.

DR EMBL; X72702; CAA51257.1; -.

DR EMBL; X80925; CAA56887.1; -.

DR EMBL; AP419525; AAL15953.1; -.

DR EMBL; AF350127; AAK57244.1; -.

DR EMBL; AF350205; AAK57322.1; -.

DR PIR; C94409; NKVLA3.

DR PIR; S32204; S32204.

DR PIR; S33686; S33686.

DR InterPro; IPR002006; Hepatitis core.

DR Pfam; PF00906; Hepatitis core; 1.

DR S32204; S32204.

DR S33686; S33686.

FT CHAIN 30 212 CORE PROTEIN

SQ SEQUENCE 212 AA; 24336 MW; 1861B13E8B047AC8 CRC64;

Query Match 99.6%; Score 790; DB 12; Length 212;

Best Local Similarity 99.3%; Pred. No. 4.1e-80;

Matches 148; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 MDIDPYKFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 60

Db 30 MDIDPYKFGATVVELLSFLPSDFPSVRDLDTASALYREALSPHCSPHTALRQAIL 89

QY 61 CWGELMTLATWGVNLEDPASRDVVSYYNTNMGLKFRQLLWFHISCLTFCGTVEYLV 120

Db 90 CWGELMTLATWGVNLEDPASRDVVSYYNTNMGLKFRQLLWFHISCLTFCGTVEYLV 149

QY 121 SFGWIRTPPAYPPNAPILSTLPETTV 149

Db 150 SFGWIRTPPAYPPNAPILSTLPETTV 178

Search completed: April 23, 2004, 16:29:38

Job time : 40 secs